SEQUENCE LISTING

<110> AMUNIX OPERATING INC.

<120> FACTOR VIII COMPOSITIONS AND METHODS OF MAKING AND USING SAME

<130> 32808-734.601

<140>
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<150> 61/599,400
<151> 2012-02-15

<160> 1700

<170> PatentIn version 3.5

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<212> PRT
<213> Homo sapiens

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Met Gln Ile Glu Leu Ser Thr Cys Phe Phe Leu Cys Leu Leu Arg Phe
1               5                  10                  15

Cys Phe Ser Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser
            20                  25                  30

Trp Asp Tyr Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg
        35                  40                  45

Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val
    50                  55                  60

Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile
65                  70                  75                  80

Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln
                85                  90                  95

Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser
            100                 105                 110

His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser
        115                 120                 125

Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp
    130                 135                 140

Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu
145                 150                 155                 160

Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser
                165                 170                 175
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Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile
            180                 185                 190

Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr
        195                 200                 205

Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly
    210                 215                 220

Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp
225                 230                 235                 240

Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr
                245                 250                 255

Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val
            260                 265                 270

Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile
        275                 280                 285

Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser
    290                 295                 300

Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met
305                 310                 315                 320

Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His
                325                 330                 335

Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro
            340                 345                 350

Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp
        355                 360                 365

Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser
    370                 375                 380

Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr
385                 390                 395                 400

Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro
                405                 410                 415

Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn
            420                 425                 430

Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met
        435                 440                 445
```

```
Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu
    450                 455                 460

Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu
465                 470                 475                 480

Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro
                485                 490                 495

His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys
            500                 505                 510

Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe
        515                 520                 525

Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp
    530                 535                 540

Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg
545                 550                 555                 560

Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu
                565                 570                 575

Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val
            580                 585                 590

Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu
        595                 600                 605

Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp
    610                 615                 620

Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val
625                 630                 635                 640

Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp
                645                 650                 655

Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe
            660                 665                 670

Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr
        675                 680                 685

Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro
    690                 695                 700

Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly
705                 710                 715                 720
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```
Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp
                725                 730                 735

Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys
            740                 745                 750

Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Ser Arg His Pro
        755                 760                 765

Ser Thr Arg Gln Lys Gln Phe Asn Ala Thr Thr Ile Pro Glu Asn Asp
    770                 775                 780

Ile Glu Lys Thr Asp Pro Trp Phe Ala His Arg Thr Pro Met Pro Lys
785                 790                 795                 800

Ile Gln Asn Val Ser Ser Ser Asp Leu Leu Met Leu Leu Arg Gln Ser
                805                 810                 815

Pro Thr Pro His Gly Leu Ser Leu Ser Asp Leu Gln Glu Ala Lys Tyr
            820                 825                 830

Glu Thr Phe Ser Asp Asp Pro Ser Pro Gly Ala Ile Asp Ser Asn Asn
        835                 840                 845

Ser Leu Ser Glu Met Thr His Phe Arg Pro Gln Leu His His Ser Gly
    850                 855                 860

Asp Met Val Phe Thr Pro Glu Ser Gly Leu Gln Leu Arg Leu Asn Glu
865                 870                 875                 880

Lys Leu Gly Thr Thr Ala Ala Thr Glu Leu Lys Lys Leu Asp Phe Lys
                885                 890                 895

Val Ser Ser Thr Ser Asn Asn Leu Ile Ser Thr Ile Pro Ser Asp Asn
            900                 905                 910

Leu Ala Ala Gly Thr Asp Asn Thr Ser Ser Leu Gly Pro Pro Ser Met
        915                 920                 925

Pro Val His Tyr Asp Ser Gln Leu Asp Thr Thr Leu Phe Gly Lys Lys
    930                 935                 940

Ser Ser Pro Leu Thr Glu Ser Gly Gly Pro Leu Ser Leu Ser Glu Glu
945                 950                 955                 960

Asn Asn Asp Ser Lys Leu Leu Glu Ser Gly Leu Met Asn Ser Gln Glu
                965                 970                 975

Ser Ser Trp Gly Lys Asn Val Ser Ser Thr Glu Ser Gly Arg Leu Phe
            980                 985                 990
```

```
Lys Gly Lys Arg Ala His Gly Pro  Ala Leu Leu Thr Lys  Asp Asn Ala
        995                 1000                1005

Leu Phe  Lys Val Ser Ile Ser  Leu Leu Lys Thr Asn  Lys Thr Ser
    1010                1015                1020

Asn Asn  Ser Ala Thr Asn Arg  Lys Thr His Ile Asp  Gly Pro Ser
    1025                1030                1035

Leu Leu  Ile Glu Asn Ser Pro  Ser Val Trp Gln Asn  Ile Leu Glu
    1040                1045                1050

Ser Asp  Thr Glu Phe Lys Lys  Val Thr Pro Leu Ile  His Asp Arg
    1055                1060                1065

Met Leu  Met Asp Lys Asn Ala  Thr Ala Leu Arg Leu  Asn His Met
    1070                1075                1080

Ser Asn  Lys Thr Thr Ser Ser  Lys Asn Met Glu Met  Val Gln Gln
    1085                1090                1095

Lys Lys  Glu Gly Pro Ile Pro  Pro Asp Ala Gln Asn  Pro Asp Met
    1100                1105                1110

Ser Phe  Phe Lys Met Leu Phe  Leu Pro Glu Ser Ala  Arg Trp Ile
    1115                1120                1125

Gln Arg  Thr His Gly Lys Asn  Ser Leu Asn Ser Gly  Gln Gly Pro
    1130                1135                1140

Ser Pro  Lys Gln Leu Val Ser  Leu Gly Pro Glu Lys  Ser Val Glu
    1145                1150                1155

Gly Gln  Asn Phe Leu Ser Glu  Lys Asn Lys Val Val  Val Gly Lys
    1160                1165                1170

Gly Glu  Phe Thr Lys Asp Val  Gly Leu Lys Glu Met  Val Phe Pro
    1175                1180                1185

Ser Ser  Arg Asn Leu Phe Leu  Thr Asn Leu Asp Asn  Leu His Glu
    1190                1195                1200

Asn Asn  Thr His Asn Gln Glu  Lys Lys Ile Gln Glu  Glu Ile Glu
    1205                1210                1215

Lys Lys  Glu Thr Leu Ile Gln  Glu Asn Val Val Leu  Pro Gln Ile
    1220                1225                1230

His Thr  Val Thr Gly Thr Lys  Asn Phe Met Lys Asn  Leu Phe Leu
    1235                1240                1245
```

```
Leu Ser  Thr Arg Gln Asn Val  Glu Gly Ser Tyr Asp  Gly Ala Tyr
    1250                1255                1260

Ala Pro  Val Leu Gln Asp Phe  Arg Ser Leu Asn Asp  Ser Thr Asn
    1265                1270                1275

Arg Thr  Lys Lys His Thr Ala  His Phe Ser Lys Lys  Gly Glu Glu
    1280                1285                1290

Glu Asn  Leu Glu Gly Leu Gly  Asn Gln Thr Lys Gln  Ile Val Glu
    1295                1300                1305

Lys Tyr  Ala Cys Thr Thr Arg  Ile Ser Pro Asn Thr  Ser Gln Gln
    1310                1315                1320

Asn Phe  Val Thr Gln Arg Ser  Lys Arg Ala Leu Lys  Gln Phe Arg
    1325                1330                1335

Leu Pro  Leu Glu Glu Thr Glu  Leu Glu Lys Arg Ile  Ile Val Asp
    1340                1345                1350

Asp Thr  Ser Thr Gln Trp Ser  Lys Asn Met Lys His  Leu Thr Pro
    1355                1360                1365

Ser Thr  Leu Thr Gln Ile Asp  Tyr Asn Glu Lys Glu  Lys Gly Ala
    1370                1375                1380

Ile Thr  Gln Ser Pro Leu Ser  Asp Cys Leu Thr Arg  Ser His Ser
    1385                1390                1395

Ile Pro  Gln Ala Asn Arg Ser  Pro Leu Pro Ile Ala  Lys Val Ser
    1400                1405                1410

Ser Phe  Pro Ser Ile Arg Pro  Ile Tyr Leu Thr Arg  Val Leu Phe
    1415                1420                1425

Gln Asp  Asn Ser Ser His Leu  Pro Ala Ala Ser Tyr  Arg Lys Lys
    1430                1435                1440

Asp Ser  Gly Val Gln Glu Ser  Ser His Phe Leu Gln  Gly Ala Lys
    1445                1450                1455

Lys Asn  Asn Leu Ser Leu Ala  Ile Leu Thr Leu Glu  Met Thr Gly
    1460                1465                1470

Asp Gln  Arg Glu Val Gly Ser  Leu Gly Thr Ser Ala  Thr Asn Ser
    1475                1480                1485

Val Thr  Tyr Lys Lys Val Glu  Asn Thr Val Leu Pro  Lys Pro Asp
    1490                1495                1500
```

```
Leu Pro  Lys Thr Ser Gly Lys  Val Glu Leu Leu Pro  Lys Val His
    1505                 1510                 1515

Ile Tyr  Gln Lys Asp Leu Phe  Pro Thr Glu Thr Ser  Asn Gly Ser
    1520                 1525                 1530

Pro Gly  His Leu Asp Leu Val  Glu Gly Ser Leu Leu  Gln Gly Thr
    1535                 1540                 1545

Glu Gly  Ala Ile Lys Trp Asn  Glu Ala Asn Arg Pro  Gly Lys Val
    1550                 1555                 1560

Pro Phe  Leu Arg Val Ala Thr  Glu Ser Ser Ala Lys  Thr Pro Ser
    1565                 1570                 1575

Lys Leu  Leu Asp Pro Leu Ala  Trp Asp Asn His Tyr  Gly Thr Gln
    1580                 1585                 1590

Ile Pro  Lys Glu Glu Trp Lys  Ser Gln Glu Lys Ser  Pro Glu Lys
    1595                 1600                 1605

Thr Ala  Phe Lys Lys Lys Asp  Thr Ile Leu Ser Leu  Asn Ala Cys
    1610                 1615                 1620

Glu Ser  Asn His Ala Ile Ala  Ala Ile Asn Glu Gly  Gln Asn Lys
    1625                 1630                 1635

Pro Glu  Ile Glu Val Thr Trp  Ala Lys Gln Gly Arg  Thr Glu Arg
    1640                 1645                 1650

Leu Cys  Ser Gln Asn Pro Pro  Val Leu Lys Arg His  Gln Arg Glu
    1655                 1660                 1665

Ile Thr  Arg Thr Thr Leu Gln  Ser Asp Gln Glu Glu  Ile Asp Tyr
    1670                 1675                 1680

Asp Asp  Thr Ile Ser Val Glu  Met Lys Lys Glu Asp  Phe Asp Ile
    1685                 1690                 1695

Tyr Asp  Glu Asp Glu Asn Gln  Ser Pro Arg Ser Phe  Gln Lys Lys
    1700                 1705                 1710

Thr Arg  His Tyr Phe Ile Ala  Ala Val Glu Arg Leu  Trp Asp Tyr
    1715                 1720                 1725

Gly Met  Ser Ser Ser Pro His  Val Leu Arg Asn Arg  Ala Gln Ser
    1730                 1735                 1740

Gly Ser  Val Pro Gln Phe Lys  Lys Val Val Phe Gln  Glu Phe Thr
    1745                 1750                 1755
```

```
Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg Gly Glu  Leu Asn Glu
    1760                1765                1770

His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg Ala Glu  Val Glu Asp
    1775                1780                1785

Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala Ser Arg  Pro Tyr Ser
    1790                1795                1800

Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu Asp Gln  Arg Gln Gly
    1805                1810                1815

Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro Asn Glu  Thr Lys Thr
    1820                1825                1830

Tyr Phe  Trp Lys Val Gln His  His Met Ala Pro Thr  Lys Asp Glu
    1835                1840                1845

Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser Asp Val  Asp Leu Glu
    1850                1855                1860

Lys Asp  Val His Ser Gly Leu  Ile Gly Pro Leu Leu  Val Cys His
    1865                1870                1875

Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg Gln Val  Thr Val Gln
    1880                1885                1890

Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp Glu Thr  Lys Ser Trp
    1895                1900                1905

Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys Arg Ala  Pro Cys Asn
    1910                1915                1920

Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu Asn Tyr  Arg Phe His
    1925                1930                1935

Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu Pro Gly  Leu Val Met
    1940                1945                1950

Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu Leu Ser  Met Gly Ser
    1955                1960                1965

Asn Glu  Asn Ile His Ser Ile  His Phe Ser Gly His  Val Phe Thr
    1970                1975                1980

Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala Leu Tyr  Asn Leu Tyr
    1985                1990                1995

Pro Gly  Val Phe Glu Thr Val  Glu Met Leu Pro Ser  Lys Ala Gly
    2000                2005                2010
```

```
Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu His Leu  His Ala Gly
    2015                 2020                 2025

Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn Lys Cys  Gln Thr Pro
    2030                 2035                 2040

Leu Gly  Met Ala Ser Gly His  Ile Arg Asp Phe Gln  Ile Thr Ala
    2045                 2050                 2055

Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys Leu Ala  Arg Leu His
    2060                 2065                 2070

Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr Lys Glu  Pro Phe Ser
    2075                 2080                 2085

Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met Ile Ile  His Gly Ile
    2090                 2095                 2100

Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser Ser Leu  Tyr Ile Ser
    2105                 2110                 2115

Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly Lys Lys  Trp Gln Thr
    2120                 2125                 2130

Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met Val Phe  Phe Gly Asn
    2135                 2140                 2145

Val Asp  Ser Ser Gly Ile Lys  His Asn Ile Phe Asn  Pro Pro Ile
    2150                 2155                 2160

Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr His Tyr  Ser Ile Arg
    2165                 2170                 2175

Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys Asp Leu  Asn Ser Cys
    2180                 2185                 2190

Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala Ile Ser  Asp Ala Gln
    2195                 2200                 2205

Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met Phe Ala  Thr Trp Ser
    2210                 2215                 2220

Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly Arg Ser  Asn Ala Trp
    2225                 2230                 2235

Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp Leu Gln  Val Asp Phe
    2240                 2245                 2250

Gln Lys  Thr Met Lys Val Thr  Gly Val Thr Thr Gln  Gly Val Lys
    2255                 2260                 2265
```

```
Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu Phe Leu  Ile Ser Ser
    2270                2275                2280

Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe Phe Gln  Asn Gly Lys
    2285                2290                2295

Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser Phe Thr  Pro Val Val
    2300                2305                2310

Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg Tyr Leu  Arg Ile His
    2315                2320                2325

Pro Gln  Ser Trp Val His Gln  Ile Ala Leu Arg Met  Glu Val Leu
    2330                2335                2340

Gly Cys  Glu Ala Gln Asp Leu  Tyr
    2345                2350
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<212> PRT
<213> Homo sapiens

<400> 2

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Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1               5                   10                  15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160
```

```
His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430
```

```
Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700
```

```
Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Ser Arg His Pro Ser Thr Arg
            740                 745                 750

Gln Lys Gln Phe Asn Ala Thr Thr Ile Pro Glu Asn Asp Ile Glu Lys
        755                 760                 765

Thr Asp Pro Trp Phe Ala His Arg Thr Pro Met Pro Lys Ile Gln Asn
    770                 775                 780

Val Ser Ser Ser Asp Leu Leu Met Leu Leu Arg Gln Ser Pro Thr Pro
785                 790                 795                 800

His Gly Leu Ser Leu Ser Asp Leu Gln Glu Ala Lys Tyr Glu Thr Phe
                805                 810                 815

Ser Asp Asp Pro Ser Pro Gly Ala Ile Asp Ser Asn Asn Ser Leu Ser
            820                 825                 830

Glu Met Thr His Phe Arg Pro Gln Leu His His Ser Gly Asp Met Val
        835                 840                 845

Phe Thr Pro Glu Ser Gly Leu Gln Leu Arg Leu Asn Glu Lys Leu Gly
    850                 855                 860

Thr Thr Ala Ala Thr Glu Leu Lys Lys Leu Asp Phe Lys Val Ser Ser
865                 870                 875                 880

Thr Ser Asn Asn Leu Ile Ser Thr Ile Pro Ser Asp Asn Leu Ala Ala
                885                 890                 895

Gly Thr Asp Asn Thr Ser Ser Leu Gly Pro Pro Ser Met Pro Val His
            900                 905                 910

Tyr Asp Ser Gln Leu Asp Thr Thr Leu Phe Gly Lys Lys Ser Ser Pro
        915                 920                 925

Leu Thr Glu Ser Gly Gly Pro Leu Ser Leu Ser Glu Glu Asn Asn Asp
    930                 935                 940

Ser Lys Leu Leu Glu Ser Gly Leu Met Asn Ser Gln Glu Ser Ser Trp
945                 950                 955                 960

Gly Lys Asn Val Ser Ser Thr Glu Ser Gly Arg Leu Phe Lys Gly Lys
                965                 970                 975
```

```
Arg Ala His Gly Pro Ala Leu Leu Thr Lys Asp Asn Ala Leu Phe Lys
            980                 985                 990

Val Ser Ile Ser Leu Leu Lys Thr  Asn Lys Thr Ser Asn  Asn Ser Ala
        995                 1000                1005

Thr Asn  Arg Lys Thr His Ile  Asp Gly Pro Ser Leu  Leu Ile Glu
    1010                1015                1020

Asn Ser  Pro Ser Val Trp Gln  Asn Ile Leu Glu Ser  Asp Thr Glu
    1025                1030                1035

Phe Lys  Lys Val Thr Pro Leu  Ile His Asp Arg Met  Leu Met Asp
    1040                1045                1050

Lys Asn  Ala Thr Ala Leu Arg  Leu Asn His Met Ser  Asn Lys Thr
    1055                1060                1065

Thr Ser  Ser Lys Asn Met Glu  Met Val Gln Gln Lys  Lys Glu Gly
    1070                1075                1080

Pro Ile  Pro Pro Asp Ala Gln  Asn Pro Asp Met Ser  Phe Phe Lys
    1085                1090                1095

Met Leu  Phe Leu Pro Glu Ser  Ala Arg Trp Ile Gln  Arg Thr His
    1100                1105                1110

Gly Lys  Asn Ser Leu Asn Ser  Gly Gln Gly Pro Ser  Pro Lys Gln
    1115                1120                1125

Leu Val  Ser Leu Gly Pro Glu  Lys Ser Val Glu Gly  Gln Asn Phe
    1130                1135                1140

Leu Ser  Glu Lys Asn Lys Val  Val Val Gly Lys Gly  Glu Phe Thr
    1145                1150                1155

Lys Asp  Val Gly Leu Lys Glu  Met Val Phe Pro Ser  Ser Arg Asn
    1160                1165                1170

Leu Phe  Leu Thr Asn Leu Asp  Asn Leu His Glu Asn  Asn Thr His
    1175                1180                1185

Asn Gln  Glu Lys Lys Ile Gln  Glu Glu Ile Glu Lys  Lys Glu Thr
    1190                1195                1200

Leu Ile  Gln Glu Asn Val Val  Leu Pro Gln Ile His  Thr Val Thr
    1205                1210                1215

Gly Thr  Lys Asn Phe Met Lys  Asn Leu Phe Leu Leu  Ser Thr Arg
    1220                1225                1230
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```
Gln Asn  Val Glu Gly Ser Tyr  Asp Gly Ala Tyr Ala  Pro Val Leu
    1235                 1240                 1245

Gln Asp  Phe Arg Ser Leu Asn  Asp Ser Thr Asn Arg  Thr Lys Lys
    1250                 1255                 1260

His Thr  Ala His Phe Ser Lys  Lys Gly Glu Glu Glu  Asn Leu Glu
    1265                 1270                 1275

Gly Leu  Gly Asn Gln Thr Lys  Gln Ile Val Glu Lys  Tyr Ala Cys
    1280                 1285                 1290

Thr Thr  Arg Ile Ser Pro Asn  Thr Ser Gln Gln Asn  Phe Val Thr
    1295                 1300                 1305

Gln Arg  Ser Lys Arg Ala Leu  Lys Gln Phe Arg Leu  Pro Leu Glu
    1310                 1315                 1320

Glu Thr  Glu Leu Glu Lys Arg  Ile Ile Val Asp Asp  Thr Ser Thr
    1325                 1330                 1335

Gln Trp  Ser Lys Asn Met Lys  His Leu Thr Pro Ser  Thr Leu Thr
    1340                 1345                 1350

Gln Ile  Asp Tyr Asn Glu Lys  Glu Lys Gly Ala Ile  Thr Gln Ser
    1355                 1360                 1365

Pro Leu  Ser Asp Cys Leu Thr  Arg Ser His Ser Ile  Pro Gln Ala
    1370                 1375                 1380

Asn Arg  Ser Pro Leu Pro Ile  Ala Lys Val Ser Ser  Phe Pro Ser
    1385                 1390                 1395

Ile Arg  Pro Ile Tyr Leu Thr  Arg Val Leu Phe Gln  Asp Asn Ser
    1400                 1405                 1410

Ser His  Leu Pro Ala Ala Ser  Tyr Arg Lys Lys Asp  Ser Gly Val
    1415                 1420                 1425

Gln Glu  Ser Ser His Phe Leu  Gln Gly Ala Lys Lys  Asn Asn Leu
    1430                 1435                 1440

Ser Leu  Ala Ile Leu Thr Leu  Glu Met Thr Gly Asp  Gln Arg Glu
    1445                 1450                 1455

Val Gly  Ser Leu Gly Thr Ser  Ala Thr Asn Ser Val  Thr Tyr Lys
    1460                 1465                 1470

Lys Val  Glu Asn Thr Val Leu  Pro Lys Pro Asp Leu  Pro Lys Thr
    1475                 1480                 1485
```

```
Ser Gly  Lys Val Glu Leu Leu  Pro Lys Val His Ile  Tyr Gln Lys
    1490                 1495                 1500

Asp Leu  Phe Pro Thr Glu Thr  Ser Asn Gly Ser Pro  Gly His Leu
    1505                 1510                 1515

Asp Leu  Val Glu Gly Ser Leu  Leu Gln Gly Thr Glu  Gly Ala Ile
    1520                 1525                 1530

Lys Trp  Asn Glu Ala Asn Arg  Pro Gly Lys Val Pro  Phe Leu Arg
    1535                 1540                 1545

Val Ala  Thr Glu Ser Ser Ala  Lys Thr Pro Ser Lys  Leu Leu Asp
    1550                 1555                 1560

Pro Leu  Ala Trp Asp Asn His  Tyr Gly Thr Gln Ile  Pro Lys Glu
    1565                 1570                 1575

Glu Trp  Lys Ser Gln Glu Lys  Ser Pro Glu Lys Thr  Ala Phe Lys
    1580                 1585                 1590

Lys Lys  Asp Thr Ile Leu Ser  Leu Asn Ala Cys Glu  Ser Asn His
    1595                 1600                 1605

Ala Ile  Ala Ala Ile Asn Glu  Gly Gln Asn Lys Pro  Glu Ile Glu
    1610                 1615                 1620

Val Thr  Trp Ala Lys Gln Gly  Arg Thr Glu Arg Leu  Cys Ser Gln
    1625                 1630                 1635

Asn Pro  Pro Val Leu Lys Arg  His Gln Arg Glu Ile  Thr Arg Thr
    1640                 1645                 1650

Thr Leu  Gln Ser Asp Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile
    1655                 1660                 1665

Ser Val  Glu Met Lys Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp
    1670                 1675                 1680

Glu Asn  Gln Ser Pro Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr
    1685                 1690                 1695

Phe Ile  Ala Ala Val Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser
    1700                 1705                 1710

Ser Pro  His Val Leu Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro
    1715                 1720                 1725

Gln Phe  Lys Lys Val Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe
    1730                 1735                 1740
```

```
Thr Gln  Pro Leu Tyr Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu
    1745                1750                1755

Leu Gly  Pro Tyr Ile Arg Ala  Glu Val Glu Asp Asn  Ile Met Val
    1760                1765                1770

Thr Phe  Arg Asn Gln Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser
    1775                1780                1785

Leu Ile  Ser Tyr Glu Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg
    1790                1795                1800

Lys Asn  Phe Val Lys Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys
    1805                1810                1815

Val Gln  His His Met Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys
    1820                1825                1830

Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu Glu Lys  Asp Val His
    1835                1840                1845

Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys His Thr  Asn Thr Leu
    1850                1855                1860

Asn Pro  Ala His Gly Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu
    1865                1870                1875

Phe Phe  Thr Ile Phe Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu
    1880                1885                1890

Asn Met  Glu Arg Asn Cys Arg  Ala Pro Cys Asn Ile  Gln Met Glu
    1895                1900                1905

Asp Pro  Thr Phe Lys Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly
    1910                1915                1920

Tyr Ile  Met Asp Thr Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln
    1925                1930                1935

Arg Ile  Arg Trp Tyr Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile
    1940                1945                1950

His Ser  Ile His Phe Ser Gly  His Val Phe Thr Val  Arg Lys Lys
    1955                1960                1965

Glu Glu  Tyr Lys Met Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe
    1970                1975                1980

Glu Thr  Val Glu Met Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val
    1985                1990                1995
```

```
Glu Cys  Leu Ile Gly Glu His  Leu His Ala Gly Met  Ser Thr Leu
    2000                 2005                 2010

Phe Leu  Val Tyr Ser Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala
    2015                 2020                 2025

Ser Gly  His Ile Arg Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr
    2030                 2035                 2040

Gly Gln  Trp Ala Pro Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser
    2045                 2050                 2055

Ile Asn  Ala Trp Ser Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val
    2060                 2065                 2070

Asp Leu  Leu Ala Pro Met Ile  Ile His Gly Ile Lys  Thr Gln Gly
    2075                 2080                 2085

Ala Arg  Gln Lys Phe Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile
    2090                 2095                 2100

Met Tyr  Ser Leu Asp Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn
    2105                 2110                 2115

Ser Thr  Gly Thr Leu Met Val  Phe Phe Gly Asn Val  Asp Ser Ser
    2120                 2125                 2130

Gly Ile  Lys His Asn Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr
    2135                 2140                 2145

Ile Arg  Leu His Pro Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg
    2150                 2155                 2160

Met Glu  Leu Met Gly Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu
    2165                 2170                 2175

Gly Met  Glu Ser Lys Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser
    2180                 2185                 2190

Ser Tyr  Phe Thr Asn Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala
    2195                 2200                 2205

Arg Leu  His Leu Gln Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val
    2210                 2215                 2220

Asn Asn  Pro Lys Glu Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met
    2225                 2230                 2235

Lys Val  Thr Gly Val Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr
    2240                 2245                 2250
```

```
Ser Met  Tyr Val Lys Glu Phe  Leu Ile Ser Ser Ser  Gln Asp Gly
    2255                 2260                 2265

His Gln  Trp Thr Leu Phe Phe  Gln Asn Gly Lys Val  Lys Val Phe
    2270                 2275                 2280

Gln Gly  Asn Gln Asp Ser Phe  Thr Pro Val Val Asn  Ser Leu Asp
    2285                 2290                 2295

Pro Pro  Leu Leu Thr Arg Tyr  Leu Arg Ile His Pro  Gln Ser Trp
    2300                 2305                 2310

Val His  Gln Ile Ala Leu Arg  Met Glu Val Leu Gly  Cys Glu Ala
    2315                 2320                 2325

Gln Asp  Leu Tyr
    2330
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Met Gln Val Glu Leu Tyr Thr Cys Cys Phe Leu Cys Leu Leu Pro Phe
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Ser Leu Ser Ala Thr Arg Lys Tyr Tyr Leu Gly Ala Val Glu Leu Ser
            20                  25                  30

Trp Asp Tyr Met Gln Ser Asp Leu Leu Ser Ala Leu His Ala Asp Thr
        35                  40                  45

Ser Phe Ser Ser Arg Val Pro Gly Ser Leu Pro Leu Thr Thr Ser Val
    50                  55                  60

Thr Tyr Arg Lys Thr Val Phe Val Glu Phe Thr Asp Asp Leu Phe Asn
65                  70                  75                  80

Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile
                85                  90                  95

Gln Ala Glu Val Tyr Asp Thr Val Val Ile Val Leu Lys Asn Met Ala
            100                 105                 110

Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala
        115                 120                 125

Ser Glu Gly Ala Glu Tyr Glu Asp Gln Thr Ser Gln Lys Glu Lys Glu
    130                 135                 140
```

```
Asp Asp Asn Val Ile Pro Gly Glu Ser His Thr Tyr Val Trp Gln Val
145                 150                 155                 160

Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Pro Cys Leu Thr Tyr
                165                 170                 175

Ser Tyr Phe Ser His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu
            180                 185                 190

Ile Gly Ala Leu Leu Val Cys Lys Glu Gly Ser Leu Ala Lys Glu Arg
        195                 200                 205

Thr Gln Thr Leu Gln Glu Phe Val Leu Leu Phe Ala Val Phe Asp Glu
    210                 215                 220

Gly Lys Ser Trp His Ser Glu Thr Asn Ala Ser Leu Thr Gln Ala Glu
225                 230                 235                 240

Ala Gln His Glu Leu His Thr Ile Asn Gly Tyr Val Asn Arg Ser Leu
                245                 250                 255

Pro Gly Leu Thr Val Cys His Lys Arg Ser Val Tyr Trp His Val Ile
            260                 265                 270

Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu Gly His
        275                 280                 285

Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile Ser Pro
    290                 295                 300

Ile Thr Phe Leu Thr Ala Gln Thr Phe Leu Met Asp Leu Gly Gln Phe
305                 310                 315                 320

Leu Leu Phe Cys His Ile Pro Ser His Gln His Asp Gly Met Glu Ala
                325                 330                 335

Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg Met Lys
            340                 345                 350

Asn Asn Glu Asp Lys Asp Tyr Asp Asp Gly Leu Tyr Asp Ser Asp Met
        355                 360                 365

Asp Val Val Ser Phe Asp Asp Asp Ser Ser Ser Pro Phe Ile Gln Ile
    370                 375                 380

Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His Tyr Ile Ala
385                 390                 395                 400

Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Ser Gly Pro Thr Pro Asn
                405                 410                 415
```

```
Asp Arg Ser His Lys Asn Leu Tyr Leu Asn Asn Gly Pro Gln Arg Ile
            420                 425                 430

Gly Lys Lys Tyr Lys Lys Val Arg Phe Val Ala Tyr Thr Asp Glu Thr
        435                 440                 445

Phe Lys Thr Arg Glu Ala Ile Gln Tyr Glu Ser Gly Ile Leu Gly Pro
    450                 455                 460

Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn
465                 470                 475                 480

Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Asn Tyr Val
                485                 490                 495

Thr Pro Leu His Thr Gly Arg Leu Pro Lys Gly Val Lys His Leu Lys
            500                 505                 510

Asp Met Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val
        515                 520                 525

Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg
    530                 535                 540

Tyr Tyr Ser Ser Phe Ile Asn Leu Glu Arg Asp Leu Ala Ser Gly Leu
545                 550                 555                 560

Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly
                565                 570                 575

Asn Gln Met Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe
            580                 585                 590

Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Met Gln Arg Phe Leu
        595                 600                 605

Pro Asn Ala Asp Val Val Gln Pro His Asp Pro Glu Phe Gln Leu Ser
    610                 615                 620

Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Asn Leu Gln Leu
625                 630                 635                 640

Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Val Gly
                645                 650                 655

Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys
            660                 665                 670

His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly
        675                 680                 685
```

```
Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Val Leu Gly
    690                 695                 700

Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys
705                 710                 715                 720

Val Ser Ser Cys Asn Arg Asn Ile Asp Asp Tyr Tyr Glu Asp Thr Tyr
                725                 730                 735

Glu Asp Ile Pro Thr Pro Leu Leu Asn Glu Asn Asn Val Ile Lys Pro
            740                 745                 750

Arg Ser Phe Ser Gln Asn Ser Arg His Pro Ser Thr Lys Glu Lys Gln
        755                 760                 765

Leu Lys Ala Thr Thr Thr Pro Glu Asn Asp Ile Glu Lys Ile Asp Leu
    770                 775                 780

Gln Ser Gly Glu Arg Thr Gln Leu Ile Lys Ala Gln Ser Val Ser Ser
785                 790                 795                 800

Ser Asp Leu Leu Met Leu Leu Gly Gln Asn Pro Thr Pro Arg Gly Leu
                805                 810                 815

Phe Leu Ser Asp Leu Arg Glu Ala Thr Asp Arg Ala Asp Asp His Ser
            820                 825                 830

Arg Gly Ala Ile Glu Arg Asn Lys Gly Pro Pro Glu Val Ala Ser Leu
        835                 840                 845

Arg Pro Glu Leu Arg His Ser Glu Asp Arg Glu Phe Thr Pro Glu Pro
    850                 855                 860

Glu Leu Gln Leu Arg Leu Asn Glu Asn Leu Gly Thr Asn Thr Thr Val
865                 870                 875                 880

Glu Leu Lys Lys Leu Asp Leu Lys Ile Ser Ser Ser Ser Asp Ser Leu
                885                 890                 895

Met Thr Ser Pro Thr Ile Pro Ser Asp Lys Leu Ala Ala Ala Thr Glu
            900                 905                 910

Lys Thr Gly Ser Leu Gly Pro Pro Asn Met Ser Val His Phe Asn Ser
        915                 920                 925

His Leu Gly Thr Ile Val Phe Gly Asn Asn Ser Ser His Leu Ile Gln
    930                 935                 940

Ser Gly Val Pro Leu Glu Leu Ser Glu Glu Asp Asn Asp Ser Lys Leu
945                 950                 955                 960
```

```
Leu Glu Ala Pro Leu Met Asn Ile Gln Glu Ser Ser Leu Arg Glu Asn
               965                 970                 975

Val Leu Ser Met Glu Ser Asn Arg Leu Phe Lys Glu Glu Arg Ile Arg
           980                 985                 990

Gly Pro Ala Ser Leu Ile Lys Asp  Asn Ala Leu Phe Lys  Val Asn Ile
       995                 1000                1005

Ser Ser  Val Lys Thr Asn Arg  Ala Pro Val Asn Leu  Thr Thr Asn
   1010                1015                1020

Arg Lys  Thr Arg Val Ala Ile  Pro Thr Leu Leu Ile  Glu Asn Ser
   1025                1030                1035

Thr Ser  Val Trp Gln Asp Ile  Met Leu Glu Arg Asn  Thr Glu Phe
   1040                1045                1050

Lys Glu  Val Thr Ser Leu Ile  His Asn Glu Thr Phe  Met Asp Arg
   1055                1060                1065

Asn Thr  Thr Ala Leu Gly Leu  Asn His Val Ser Asn  Lys Thr Thr
   1070                1075                1080

Leu Ser  Lys Asn Val Glu Met  Ala His Gln Lys Lys  Glu Asp Pro
   1085                1090                1095

Val Pro  Leu Arg Ala Glu Asn  Pro Asp Leu Ser Ser  Ser Lys Ile
   1100                1105                1110

Pro Phe  Leu Pro Asp Trp Ile  Lys Thr His Gly Lys  Asn Ser Leu
   1115                1120                1125

Ser Ser  Glu Gln Arg Pro Ser  Pro Lys Gln Leu Thr  Ser Leu Gly
   1130                1135                1140

Ser Glu  Lys Ser Val Lys Asp  Gln Asn Phe Leu Ser  Glu Glu Lys
   1145                1150                1155

Val Val  Val Gly Glu Asp Glu  Phe Thr Lys Asp Thr  Glu Leu Gln
   1160                1165                1170

Glu Ile  Phe Pro Asn Asn Lys  Ser Ile Phe Phe Ala  Asn Leu Ala
   1175                1180                1185

Asn Val  Gln Glu Asn Asp Thr  Tyr Asn Gln Glu Lys  Lys Ser Pro
   1190                1195                1200

Glu Glu  Ile Glu Arg Lys Glu  Lys Leu Thr Gln Glu  Asn Val Ala
   1205                1210                1215
```

```
Leu Pro  Gln Ala His Thr Met  Ile Gly Thr Lys Asn  Phe Leu Lys
    1220                1225                1230

Asn Leu  Phe Leu Leu Ser Thr  Lys Gln Asn Val Ala  Gly Leu Glu
    1235                1240                1245

Glu Gln  Pro Tyr Thr Pro Ile  Leu Gln Asp Thr Arg  Ser Leu Asn
    1250                1255                1260

Asp Ser  Pro His Ser Glu Gly  Ile His Met Ala Asn  Phe Ser Lys
    1265                1270                1275

Ile Arg  Glu Glu Ala Asn Leu  Glu Gly Leu Gly Asn  Gln Thr Asn
    1280                1285                1290

Gln Met  Val Glu Arg Phe Pro  Ser Thr Thr Arg Met  Ser Ser Asn
    1295                1300                1305

Ala Ser  Gln His Val Ile Thr  Gln Arg Gly Lys Arg  Ser Leu Lys
    1310                1315                1320

Gln Pro  Arg Leu Ser Gln Gly  Glu Ile Lys Phe Glu  Arg Lys Val
    1325                1330                1335

Ile Ala  Asn Asp Thr Ser Thr  Gln Trp Ser Lys Asn  Met Asn Tyr
    1340                1345                1350

Leu Ala  Gln Gly Thr Leu Thr  Gln Ile Glu Tyr Asn  Glu Lys Glu
    1355                1360                1365

Lys Arg  Ala Ile Thr Gln Ser  Pro Leu Ser Asp Cys  Ser Met Arg
    1370                1375                1380

Asn His  Val Thr Ile Gln Met  Asn Asp Ser Ala Leu  Pro Val Ala
    1385                1390                1395

Lys Glu  Ser Ala Ser Pro Ser  Val Arg His Thr Asp  Leu Thr Lys
    1400                1405                1410

Ile Pro  Ser Gln His Asn Ser  Ser His Leu Pro Ala  Ser Ala Cys
    1415                1420                1425

Asn Tyr  Thr Phe Arg Glu Arg  Thr Ser Gly Val Gln  Glu Gly Ser
    1430                1435                1440

His Phe  Leu Gln Glu Ala Lys  Arg Asn Asn Leu Ser  Leu Ala Phe
    1445                1450                1455

Val Thr  Leu Gly Ile Thr Glu  Gly Gln Gly Lys Phe  Ser Ser Leu
    1460                1465                1470
```

```
Gly Lys  Ser Ala Thr Asn Gln  Pro Met Tyr Lys Lys  Leu Glu Asn
    1475                1480                1485

Thr Val  Leu Leu Gln Pro Gly  Leu Ser Glu Thr Ser  Asp Lys Val
    1490                1495                1500

Glu Leu  Leu Ser Gln Val His  Val Asp Gln Glu Asp  Ser Phe Pro
    1505                1510                1515

Thr Lys  Thr Ser Asn Asp Ser  Pro Gly His Leu Asp  Leu Met Gly
    1520                1525                1530

Lys Ile  Phe Leu Gln Lys Thr  Gln Gly Pro Val Lys  Met Asn Lys
    1535                1540                1545

Thr Asn  Ser Pro Gly Lys Val  Pro Phe Leu Lys Trp  Ala Thr Glu
    1550                1555                1560

Ser Ser  Glu Lys Ile Pro Ser  Lys Leu Leu Gly Val  Leu Ala Trp
    1565                1570                1575

Asp Asn  His Tyr Asp Thr Gln  Ile Pro Ser Glu Glu  Trp Lys Ser
    1580                1585                1590

Gln Lys  Lys Ser Gln Thr Asn  Thr Ala Phe Lys Arg  Lys Asp Thr
    1595                1600                1605

Ile Leu  Pro Leu Gly Pro Cys  Glu Asn Asn Asp Ser  Thr Ala Ala
    1610                1615                1620

Ile Asn  Glu Gly Gln Asp Lys  Pro Gln Arg Glu Ala  Met Trp Ala
    1625                1630                1635

Lys Gln  Gly Glu Pro Gly Arg  Leu Cys Ser Gln Asn  Pro Pro Val
    1640                1645                1650

Ser Lys  His His Gln Arg Glu  Ile Thr Val Thr Thr  Leu Gln Pro
    1655                1660                1665

Glu Glu  Asp Lys Phe Glu Tyr  Asp Asp Thr Phe Ser  Ile Glu Met
    1670                1675                1680

Lys Arg  Glu Asp Phe Asp Ile  Tyr Gly Asp Tyr Glu  Asn Gln Gly
    1685                1690                1695

Leu Arg  Ser Phe Gln Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala
    1700                1705                1710

Val Glu  Arg Leu Trp Asp Tyr  Gly Met Ser Arg Ser  Pro His Ile
    1715                1720                1725
```

```
Leu Arg  Asn Arg Ala Gln Ser  Gly Asp Val Gln Gln  Phe Lys Lys
    1730                1735                1740

Val Val  Phe Gln Glu Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu
    1745                1750                1755

Tyr Arg  Gly Glu Leu Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr
    1760                1765                1770

Ile Arg  Ala Glu Val Glu Asp  Asn Ile Val Val Thr  Phe Lys Asn
    1775                1780                1785

Gln Ala  Ser Arg Pro Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr
    1790                1795                1800

Asp Glu  Asp Glu Gly Gln Gly  Ala Glu Pro Arg Arg  Lys Phe Val
    1805                1810                1815

Asn Pro  Asn Glu Thr Lys Ile  Tyr Phe Trp Lys Val  Gln His His
    1820                1825                1830

Met Ala  Pro Thr Lys Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr
    1835                1840                1845

Phe Ser  Asp Val Asp Leu Glu  Lys Asp Val His Ser  Gly Leu Ile
    1850                1855                1860

Gly Pro  Leu Leu Ile Cys Arg  Ser Asn Thr Leu Asn  Pro Ala His
    1865                1870                1875

Gly Arg  Gln Val Thr Val Gln  Glu Phe Ala Leu Val  Phe Thr Ile
    1880                1885                1890

Phe Asp  Glu Thr Lys Ser Trp  Tyr Phe Thr Glu Asn  Leu Glu Arg
    1895                1900                1905

Asn Cys  Arg Ala Pro Cys Asn  Val Gln Lys Glu Asp  Pro Thr Leu
    1910                1915                1920

Lys Glu  Asn Phe Arg Phe His  Ala Ile Asn Gly Tyr  Val Lys Asp
    1925                1930                1935

Thr Leu  Pro Gly Leu Val Met  Ala Gln Asp Gln Lys  Val Arg Trp
    1940                1945                1950

Tyr Leu  Leu Ser Met Gly Ser  Asn Glu Asn Ile His  Ser Ile His
    1955                1960                1965

Phe Ser  Gly His Val Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys
    1970                1975                1980
```

```
Met Ala  Val Tyr Asn Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu
    1985                 1990                 1995

Met Leu  Pro Ser Gln Val Gly  Ile Trp Arg Ile Glu  Cys Leu Ile
    2000                 2005                 2010

Gly Glu  His Leu Gln Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr
    2015                 2020                 2025

Ser Lys  Lys Cys Gln Thr Pro  Leu Gly Met Ala Ser  Gly His Ile
    2030                 2035                 2040

Arg Asp  Phe Gln Ile Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala
    2045                 2050                 2055

Pro Lys  Leu Ala Arg Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp
    2060                 2065                 2070

Ser Thr  Lys Asp Pro Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala
    2075                 2080                 2085

Pro Met  Ile Ile His Gly Ile  Met Thr Gln Gly Ala  Arg Gln Lys
    2090                 2095                 2100

Phe Ser  Ser Leu Tyr Val Ser  Gln Phe Ile Ile Met  Tyr Ser Leu
    2105                 2110                 2115

Asp Gly  Asn Lys Trp His Ser  Tyr Arg Gly Asn Ser  Thr Gly Thr
    2120                 2125                 2130

Leu Met  Val Phe Phe Gly Asn  Val Asp Ser Ser Gly  Ile Lys His
    2135                 2140                 2145

Asn Ile  Phe Asn Pro Pro Ile  Ile Ala Gln Tyr Ile  Arg Leu His
    2150                 2155                 2160

Pro Thr  His Tyr Ser Ile Arg  Ser Thr Leu Arg Met  Glu Leu Leu
    2165                 2170                 2175

Gly Cys  Asp Phe Asn Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser
    2180                 2185                 2190

Lys Ala  Ile Ser Asp Ala Gln  Ile Thr Ala Ser Ser  Tyr Leu Ser
    2195                 2200                 2205

Ser Met  Leu Ala Thr Trp Ser  Pro Ser Gln Ala Arg  Leu His Leu
    2210                 2215                 2220

Gln Gly  Arg Thr Asn Ala Trp  Arg Pro Gln Ala Asn  Asn Pro Lys
    2225                 2230                 2235
```

Glu Trp Leu Gln Val Asp Phe Arg Lys Thr Met Lys Val Thr Gly
2240 2245 2250

Ile Thr Thr Gln Gly Val Lys Ser Leu Leu Ile Ser Met Tyr Val
2255 2260 2265

Lys Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly His Asn Trp Thr
2270 2275 2280

Leu Phe Leu Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn Arg
2285 2290 2295

Asp Ser Ser Thr Pro Val Arg Asn Arg Leu Glu Pro Pro Leu Val
2300 2305 2310

Ala Arg Tyr Val Arg Leu His Pro Gln Ser Trp Ala His His Ile
2315 2320 2325

Ala Leu Arg Leu Glu Val Leu Gly Cys Asp Thr Gln Gln Pro Ala
2330 2335 2340

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

```
Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415
```

```
Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685
```

```
Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Ser Arg His Pro Ser Thr Arg
            740                 745                 750

Gln Lys Gln Phe Asn Ala Thr Thr Ile Pro Glu Asn Asp Ile Glu Lys
        755                 760                 765

Thr Asp Pro Trp Phe Ala His Arg Thr Pro Met Pro Lys Ile Gln Asn
    770                 775                 780

Val Ser Ser Ser Asp Leu Leu Met Leu Leu Arg Gln Ser Pro Thr Pro
785                 790                 795                 800

His Gly Leu Ser Leu Ser Asp Leu Gln Glu Ala Lys Tyr Glu Thr Phe
                805                 810                 815

Ser Asp Asp Pro Ser Pro Gly Ala Ile Asp Ser Asn Asn Ser Leu Ser
            820                 825                 830

Glu Met Thr His Phe Arg Pro Gln Leu His His Ser Gly Asp Met Val
        835                 840                 845

Phe Thr Pro Glu Ser Gly Leu Gln Leu Arg Leu Asn Glu Lys Leu Gly
    850                 855                 860

Thr Thr Ala Ala Thr Glu Leu Lys Lys Leu Asp Phe Lys Val Ser Ser
865                 870                 875                 880

Thr Ser Asn Asn Leu Ile Ser Thr Ile Pro Ser Asp Asn Leu Ala Ala
                885                 890                 895

Gly Thr Asp Asn Thr Ser Ser Leu Gly Pro Pro Ser Met Pro Val His
            900                 905                 910

Tyr Asp Ser Gln Leu Asp Thr Thr Leu Phe Gly Lys Lys Ser Ser Pro
        915                 920                 925

Leu Thr Glu Ser Gly Gly Pro Leu Ser Leu Ser Glu Glu Asn Asn Asp
    930                 935                 940

Ser Lys Leu Leu Glu Ser Gly Leu Met Asn Ser Gln Glu Ser Ser Trp
945                 950                 955                 960
```

```
Gly Lys Asn Val Ser Ser Thr Glu Ser Gly Arg Leu Phe Lys Gly Lys
                965                 970                 975

Arg Ala His Gly Pro Ala Leu Leu Thr Lys Asp Asn Ala Leu Phe Lys
            980                 985                 990

Val Ser Ile Ser Leu Leu Lys Thr  Asn Lys Thr Ser Asn  Asn Ser Ala
        995                 1000                1005

Thr Asn  Arg Lys Thr His Ile  Asp Gly Pro Ser Leu  Leu Ile Glu
    1010                1015                1020

Asn Ser  Pro Ser Val Trp Gln  Asn Ile Leu Glu Ser  Asp Thr Glu
    1025                1030                1035

Phe Lys  Lys Val Thr Pro Leu  Ile His Asp Arg Met  Leu Met Asp
    1040                1045                1050

Lys Asn  Ala Thr Ala Leu Arg  Leu Asn His Met Ser  Asn Lys Thr
    1055                1060                1065

Thr Ser  Ser Lys Asn Met Glu  Met Val Gln Gln Lys  Lys Glu Gly
    1070                1075                1080

Pro Ile  Pro Pro Asp Ala Gln  Asn Pro Asp Met Ser  Phe Phe Lys
    1085                1090                1095

Met Leu  Phe Leu Pro Glu Ser  Ala Arg Trp Ile Gln  Arg Thr His
    1100                1105                1110

Gly Lys  Asn Ser Leu Asn Ser  Gly Gln Gly Pro Ser  Pro Lys Gln
    1115                1120                1125

Leu Val  Ser Leu Gly Pro Glu  Lys Ser Val Glu Gly  Gln Asn Phe
    1130                1135                1140

Leu Ser  Glu Lys Asn Lys Val  Val Val Gly Lys Gly  Glu Phe Thr
    1145                1150                1155

Lys Asp  Val Gly Leu Lys Glu  Met Val Phe Pro Ser  Ser Arg Asn
    1160                1165                1170

Leu Phe  Leu Thr Asn Leu Asp  Asn Leu His Glu Asn  Asn Thr His
    1175                1180                1185

Asn Gln  Glu Lys Lys Ile Gln  Glu Glu Ile Glu Lys  Lys Glu Thr
    1190                1195                1200

Leu Ile  Gln Glu Asn Val Val  Leu Pro Gln Ile His  Thr Val Thr
    1205                1210                1215
```

```
Gly Thr  Lys Asn Phe Met Lys  Asn Leu Phe Leu Leu  Ser Thr Arg
    1220                 1225                 1230

Gln Asn  Val Glu Gly Ser Tyr  Asp Gly Ala Tyr Ala  Pro Val Leu
    1235                 1240                 1245

Gln Asp  Phe Arg Ser Leu Asn  Asp Ser Thr Asn Arg  Thr Lys Lys
    1250                 1255                 1260

His Thr  Ala His Phe Ser Lys  Lys Gly Glu Glu Glu  Asn Leu Glu
    1265                 1270                 1275

Gly Leu  Gly Asn Gln Thr Lys  Gln Ile Val Glu Lys  Tyr Ala Cys
    1280                 1285                 1290

Thr Thr  Arg Ile Ser Pro Asn  Thr Ser Gln Gln Asn  Phe Val Thr
    1295                 1300                 1305

Gln Arg  Ser Lys Arg Ala Leu  Lys Gln Phe Arg Leu  Pro Leu Glu
    1310                 1315                 1320

Glu Thr  Glu Leu Glu Lys Arg  Ile Ile Val Asp Asp  Thr Ser Thr
    1325                 1330                 1335

Gln Trp  Ser Lys Asn Met Lys  His Leu Thr Pro Ser  Thr Leu Thr
    1340                 1345                 1350

Gln Ile  Asp Tyr Asn Glu Lys  Glu Lys Gly Ala Ile  Thr Gln Ser
    1355                 1360                 1365

Pro Leu  Ser Asp Cys Leu Thr  Arg Ser His Ser Ile  Pro Gln Ala
    1370                 1375                 1380

Asn Arg  Ser Pro Leu Pro Ile  Ala Lys Val Ser Ser  Phe Pro Ser
    1385                 1390                 1395

Ile Arg  Pro Ile Tyr Leu Thr  Arg Val Leu Phe Gln  Asp Asn Ser
    1400                 1405                 1410

Ser His  Leu Pro Ala Ala Ser  Tyr Arg Lys Lys Asp  Ser Gly Val
    1415                 1420                 1425

Gln Glu  Ser Ser His Phe Leu  Gln Gly Ala Lys Lys  Asn Asn Leu
    1430                 1435                 1440

Ser Leu  Ala Ile Leu Thr Leu  Glu Met Thr Gly Asp  Gln Arg Glu
    1445                 1450                 1455

Val Gly  Ser Leu Gly Thr Ser  Ala Thr Asn Ser Val  Thr Tyr Lys
    1460                 1465                 1470
```

```
Lys Val  Glu Asn Thr Val Leu  Pro Lys Pro Asp Leu  Pro Lys Thr
    1475                 1480                 1485

Ser Gly  Lys Val Glu Leu Leu  Pro Lys Val His Ile  Tyr Gln Lys
    1490                 1495                 1500

Asp Leu  Phe Pro Thr Glu Thr  Ser Asn Gly Ser Pro  Gly His Leu
    1505                 1510                 1515

Asp Leu  Val Glu Gly Ser Leu  Leu Gln Gly Thr Glu  Gly Ala Ile
    1520                 1525                 1530

Lys Trp  Asn Glu Ala Asn Arg  Pro Gly Lys Val Pro  Phe Leu Arg
    1535                 1540                 1545

Val Ala  Thr Glu Ser Ser Ala  Lys Thr Pro Ser Lys  Leu Leu Asp
    1550                 1555                 1560

Pro Leu  Ala Trp Asp Asn His  Tyr Gly Thr Gln Ile  Pro Lys Glu
    1565                 1570                 1575

Glu Trp  Lys Ser Gln Glu Lys  Ser Pro Glu Lys Thr  Ala Phe Lys
    1580                 1585                 1590

Lys Lys  Asp Thr Ile Leu Ser  Leu Asn Ala Cys Glu  Ser Asn His
    1595                 1600                 1605

Ala Ile  Ala Ala Ile Asn Glu  Gly Gln Asn Lys Pro  Glu Ile Glu
    1610                 1615                 1620

Val Thr  Trp Ala Lys Gln Gly  Arg Thr Glu Arg Leu  Cys Ser Gln
    1625                 1630                 1635

Asn Pro  Pro Val Leu Lys Arg  His Gln Arg Glu Ile  Thr Arg Thr
    1640                 1645                 1650

Thr Leu  Gln Ser Asp Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile
    1655                 1660                 1665

Ser Val  Glu Met Lys Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp
    1670                 1675                 1680

Glu Asn  Gln Ser Pro Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr
    1685                 1690                 1695

Phe Ile  Ala Ala Val Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser
    1700                 1705                 1710

Ser Pro  His Val Leu Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro
    1715                 1720                 1725
```

```
Gln Phe  Lys Lys Val Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe
    1730                1735                1740

Thr Gln  Pro Leu Tyr Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu
    1745                1750                1755

Leu Gly  Pro Tyr Ile Arg Ala  Glu Val Glu Asp Asn  Ile Met Val
    1760                1765                1770

Thr Phe  Arg Asn Gln Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser
    1775                1780                1785

Leu Ile  Ser Tyr Glu Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg
    1790                1795                1800

Lys Asn  Phe Val Lys Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys
    1805                1810                1815

Val Gln  His His Met Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys
    1820                1825                1830

Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu Glu Lys  Asp Val His
    1835                1840                1845

Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys His Thr  Asn Thr Leu
    1850                1855                1860

Asn Pro  Ala His Gly Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu
    1865                1870                1875

Phe Phe  Thr Ile Phe Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu
    1880                1885                1890

Asn Met  Glu Arg Asn Cys Arg  Ala Pro Cys Asn Ile  Gln Met Glu
    1895                1900                1905

Asp Pro  Thr Phe Lys Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly
    1910                1915                1920

Tyr Ile  Met Asp Thr Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln
    1925                1930                1935

Arg Ile  Arg Trp Tyr Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile
    1940                1945                1950

His Ser  Ile His Phe Ser Gly  His Val Phe Thr Val  Arg Lys Lys
    1955                1960                1965

Glu Glu  Tyr Lys Met Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe
    1970                1975                1980
```

```
Glu Thr  Val Glu Met Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val
    1985                 1990                 1995

Glu Cys  Leu Ile Gly Glu His  Leu His Ala Gly Met  Ser Thr Leu
    2000                 2005                 2010

Phe Leu  Val Tyr Ser Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala
    2015                 2020                 2025

Ser Gly  His Ile Arg Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr
    2030                 2035                 2040

Gly Gln  Trp Ala Pro Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser
    2045                 2050                 2055

Ile Asn  Ala Trp Ser Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val
    2060                 2065                 2070

Asp Leu  Leu Ala Pro Met Ile  Ile His Gly Ile Lys  Thr Gln Gly
    2075                 2080                 2085

Ala Arg  Gln Lys Phe Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile
    2090                 2095                 2100

Met Tyr  Ser Leu Asp Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn
    2105                 2110                 2115

Ser Thr  Gly Thr Leu Met Val  Phe Phe Gly Asn Val  Asp Ser Ser
    2120                 2125                 2130

Gly Ile  Lys His Asn Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr
    2135                 2140                 2145

Ile Arg  Leu His Pro Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg
    2150                 2155                 2160

Met Glu  Leu Met Gly Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu
    2165                 2170                 2175

Gly Met  Glu Ser Lys Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser
    2180                 2185                 2190

Ser Tyr  Phe Thr Asn Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala
    2195                 2200                 2205

Arg Leu  His Leu Gln Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val
    2210                 2215                 2220

Asn Asn  Pro Lys Glu Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met
    2225                 2230                 2235
```

```
Lys Val  Thr Gly Val Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr
    2240                 2245                 2250

Ser Met  Tyr Val Lys Glu Phe  Leu Ile Ser Ser Ser  Gln Asp Gly
    2255                 2260                 2265

His Gln  Trp Thr Leu Phe Phe  Gln Asn Gly Lys Val  Lys Val Phe
    2270                 2275                 2280

Gln Gly  Asn Gln Asp Ser Phe  Thr Pro Val Val Asn  Ser Leu Asp
    2285                 2290                 2295

Pro Pro  Leu Leu Thr Arg Tyr  Leu Arg Ile His Pro  Gln Ser Trp
    2300                 2305                 2310

Val His  Gln Ile Ala Leu Arg  Met Glu Val Leu Gly  Cys Glu Ala
    2315                 2320                 2325

Gln Asp  Leu Tyr
    2330
```

<210> 5
<211> 2300
<212> PRT
<213> Mus sp.

<400> 5

```
Ala Ile Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asn Tyr
1               5                  10                  15

Ile Gln Ser Asp Leu Leu Ser Val Leu His Thr Asp Ser Arg Phe Leu
            20                  25                  30

Pro Arg Met Ser Thr Ser Phe Pro Phe Asn Thr Ser Ile Met Tyr Lys
        35                  40                  45

Lys Thr Val Phe Val Glu Tyr Lys Asp Gln Leu Phe Asn Ile Ala Lys
    50                  55                  60

Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Trp Thr Glu
65                  70                  75                  80

Val His Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro
                85                  90                  95

Val Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly
            100                 105                 110

Asp Glu Tyr Glu Asp Gln Thr Ser Gln Met Glu Lys Glu Asp Asp Lys
        115                 120                 125
```

```
Val Phe Pro Gly Glu Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu
    130                 135                 140

Asn Gly Pro Met Ala Ser Asp Pro Pro Cys Leu Thr Tyr Ser Tyr Met
145                 150                 155                 160

Ser His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala
                165                 170                 175

Leu Leu Val Cys Lys Glu Gly Ser Leu Ser Lys Glu Arg Thr Gln Met
            180                 185                 190

Leu Tyr Gln Phe Val Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser
        195                 200                 205

Trp His Ser Glu Thr Asn Asp Ser Tyr Thr Gln Ser Met Asp Ser Ala
    210                 215                 220

Ser Ala Arg Asp Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn
225                 230                 235                 240

Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp
                245                 250                 255

His Val Ile Gly Met Gly Thr Thr Pro Glu Ile His Ser Ile Phe Leu
            260                 265                 270

Glu Gly His Thr Phe Phe Val Arg Asn His Arg Gln Ala Ser Leu Glu
        275                 280                 285

Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Ile Asp Leu
    290                 295                 300

Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Lys His Asp Gly
305                 310                 315                 320

Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Ser Gln Trp
                325                 330                 335

Gln Lys Lys Asn Asn Asn Glu Glu Met Glu Asp Tyr Asp Asp Asp Leu
            340                 345                 350

Tyr Ser Glu Met Asp Met Phe Thr Leu Asp Tyr Asp Ser Ser Pro Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys Tyr Pro Lys Thr Trp Ile His
    370                 375                 380

Tyr Ile Ser Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Ser Val Pro
385                 390                 395                 400
```

```
Thr Ser Asp Asn Gly Ser Tyr Lys Ser Gln Tyr Leu Ser Asn Gly Pro
                405                 410                 415

His Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Ile Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Thr Ile Gln His Glu Ser Gly Leu
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Ser Pro Leu His Ala Arg Arg Leu Pro Arg Gly Ile Lys
                485                 490                 495

His Val Lys Asp Leu Pro Ile His Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Ile Asn Pro Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Met Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Ile Phe Asp Glu Asn Gln Ser Trp Tyr Ile Thr Glu Asn Met Gln
            580                 585                 590

Arg Phe Leu Pro Asn Ala Ala Lys Thr Gln Pro Gln Asp Pro Gly Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Glu Leu Thr Val Cys Leu His Glu Val Ala Tyr Trp His Ile Leu
625                 630                 635                 640

Ser Val Gly Ala Gln Thr Asp Phe Leu Ser Ile Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670
```

```
Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Val Leu Gly Cys His Asn Ser Asp Phe Arg Lys Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Ser Thr Ser Asp Tyr Tyr Glu
705                 710                 715                 720

Glu Ile Tyr Glu Asp Ile Pro Thr Gln Leu Val Asn Glu Asn Asn Val
                725                 730                 735

Ile Asp Pro Arg Ser Phe Phe Gln Asn Thr Asn His Pro Asn Thr Arg
            740                 745                 750

Lys Lys Lys Phe Lys Asp Ser Thr Ile Pro Lys Asn Asp Met Glu Lys
        755                 760                 765

Ile Glu Pro Gln Phe Glu Glu Ile Ala Glu Met Leu Lys Val Gln Ser
    770                 775                 780

Val Ser Val Ser Asp Met Leu Met Leu Leu Gly Gln Ser His Pro Thr
785                 790                 795                 800

Pro His Gly Leu Phe Leu Ser Asp Gly Gln Glu Ala Ile Tyr Glu Ala
                805                 810                 815

Ile His Asp Asp His Ser Pro Asn Ala Ile Asp Ser Asn Glu Gly Pro
            820                 825                 830

Ser Lys Val Thr Gln Leu Arg Pro Glu Ser His His Ser Glu Lys Ile
        835                 840                 845

Val Phe Thr Pro Gln Pro Gly Leu Gln Leu Arg Ser Asn Lys Ser Leu
    850                 855                 860

Glu Thr Thr Ile Glu Val Lys Trp Lys Lys Leu Gly Leu Gln Val Ser
865                 870                 875                 880

Ser Leu Pro Ser Asn Leu Met Thr Thr Thr Ile Leu Ser Asp Asn Leu
                885                 890                 895

Lys Ala Thr Phe Glu Lys Thr Asp Ser Ser Gly Phe Pro Asp Met Pro
            900                 905                 910

Val His Ser Ser Ser Lys Leu Ser Thr Thr Ala Phe Gly Lys Lys Ala
        915                 920                 925

Tyr Ser Leu Val Gly Ser His Val Pro Leu Asn Ala Ser Glu Glu Asn
    930                 935                 940
```

```
Ser Asp Ser Asn Ile Leu Asp Ser Thr Leu Met Tyr Ser Gln Glu Ser
945                 950                 955                 960

Leu Pro Arg Asp Asn Ile Leu Ser Ile Glu Asn Asp Arg Leu Leu Arg
                965                 970                 975

Glu Lys Arg Phe His Gly Ile Ala Leu Leu Thr Lys Asp Asn Thr Leu
            980                 985                 990

Phe Lys Asp Asn Val Ser Leu Met  Lys Thr Asn Lys Thr  Tyr Asn His
        995                 1000                1005

Ser Thr  Thr Asn Glu Lys Leu  His Thr Glu Ser Pro  Thr Ser Ile
    1010                1015                1020

Glu Asn  Ser Thr Thr Asp Leu  Gln Asp Ala Ile Leu  Lys Val Asn
    1025                1030                1035

Ser Glu  Ile Gln Glu Val Thr  Ala Leu Ile His Asp  Gly Thr Leu
    1040                1045                1050

Leu Gly  Lys Asn Ser Thr Tyr  Leu Arg Leu Asn His  Met Leu Asn
    1055                1060                1065

Arg Thr  Thr Ser Thr Lys Asn  Lys Asp Ile Phe His  Arg Lys Asp
    1070                1075                1080

Glu Asp  Pro Ile Pro Gln Asp  Glu Glu Asn Thr Ile  Met Pro Phe
    1085                1090                1095

Ser Lys  Met Leu Phe Leu Ser  Glu Ser Ser Asn Trp  Phe Lys Lys
    1100                1105                1110

Thr Asn  Gly Asn Asn Ser Leu  Asn Ser Glu Gln Glu  His Ser Pro
    1115                1120                1125

Lys Gln  Leu Val Tyr Leu Met  Phe Lys Lys Tyr Val  Lys Asn Gln
    1130                1135                1140

Ser Phe  Leu Ser Glu Lys Asn  Lys Val Thr Val Glu  Gln Asp Gly
    1145                1150                1155

Phe Thr  Lys Asn Ile Gly Leu  Lys Asp Met Ala Phe  Pro His Asn
    1160                1165                1170

Met Ser  Ile Phe Leu Thr Thr  Leu Ser Asn Val His  Glu Asn Gly
    1175                1180                1185

Arg His  Asn Gln Glu Lys Asn  Ile Gln Glu Glu Ile  Glu Lys Glu
    1190                1195                1200
```

```
Ala Leu  Ile Glu Glu Lys Val  Val Leu Pro Gln Val  His Glu Ala
    1205                 1210                 1215

Thr Gly  Ser Lys Asn Phe Leu  Lys Asp Ile Leu Ile  Leu Gly Thr
    1220                 1225                 1230

Arg Gln  Asn Ile Ser Leu Tyr  Glu Val His Val Pro  Val Leu Gln
    1235                 1240                 1245

Asn Ile  Thr Ser Ile Asn Asn  Ser Thr Asn Thr Val  Gln Ile His
    1250                 1255                 1260

Met Glu  His Phe Phe Lys Arg  Arg Lys Asp Lys Glu  Thr Asn Ser
    1265                 1270                 1275

Glu Gly  Leu Val Asn Lys Thr  Arg Glu Met Val Lys  Asn Tyr Pro
    1280                 1285                 1290

Ser Gln  Lys Asn Ile Thr Thr  Gln Arg Ser Lys Arg  Ala Leu Gly
    1295                 1300                 1305

Gln Phe  Arg Leu Ser Thr Gln  Trp Leu Lys Thr Ile  Asn Cys Ser
    1310                 1315                 1320

Thr Gln  Cys Ile Ile Lys Gln  Ile Asp His Ser Lys  Glu Met Lys
    1325                 1330                 1335

Lys Phe  Ile Thr Lys Ser Ser  Leu Ser Asp Ser Ser  Val Ile Lys
    1340                 1345                 1350

Ser Thr  Thr Gln Thr Asn Ser  Ser Asp Ser His Ile  Val Lys Thr
    1355                 1360                 1365

Ser Ala  Phe Pro Pro Ile Asp  Leu Lys Arg Ser Pro  Phe Gln Asn
    1370                 1375                 1380

Lys Phe  Ser His Val Gln Ala  Ser Ser Tyr Ile Tyr  Asp Phe Lys
    1385                 1390                 1395

Thr Lys  Ser Ser Arg Ile Gln  Glu Ser Asn Asn Phe  Leu Lys Glu
    1400                 1405                 1410

Thr Lys  Ile Asn Asn Pro Ser  Leu Ala Ile Leu Pro  Trp Asn Met
    1415                 1420                 1425

Phe Ile  Asp Gln Gly Lys Phe  Thr Ser Pro Gly Lys  Ser Asn Thr
    1430                 1435                 1440

Asn Ser  Val Thr Tyr Lys Lys  Arg Glu Asn Ile Ile  Phe Leu Lys
    1445                 1450                 1455
```

```
Pro Thr  Leu Pro Glu Glu Ser  Gly Lys Ile Glu Leu  Leu Pro Gln
    1460                1465                1470

Val Ser  Ile Gln Glu Glu Glu  Ile Leu Pro Thr Glu  Thr Ser His
    1475                1480                1485

Gly Ser  Pro Gly His Leu Asn  Leu Met Lys Glu Val  Phe Leu Gln
    1490                1495                1500

Lys Ile  Gln Gly Pro Thr Lys  Trp Asn Lys Ala Lys  Arg His Gly
    1505                1510                1515

Glu Ser  Ile Lys Gly Lys Thr  Glu Ser Ser Lys Asn  Thr Arg Ser
    1520                1525                1530

Lys Leu  Leu Asn His His Ala  Trp Asp Tyr His Tyr  Ala Ala Gln
    1535                1540                1545

Ile Pro  Lys Asp Met Trp Lys  Ser Lys Glu Lys Ser  Pro Glu Ile
    1550                1555                1560

Ile Ser  Ile Lys Gln Glu Asp  Thr Ile Leu Ser Leu  Arg Pro His
    1565                1570                1575

Gly Asn  Ser His Ser Ile Gly  Ala Asn Glu Lys Gln  Asn Trp Pro
    1580                1585                1590

Gln Arg  Glu Thr Thr Trp Val  Lys Gln Gly Gln Thr  Gln Arg Thr
    1595                1600                1605

Cys Ser  Gln Ile Pro Pro Val  Leu Lys Arg His Gln  Arg Glu Leu
    1610                1615                1620

Ser Ala  Phe Gln Ser Glu Gln  Glu Ala Thr Asp Tyr  Asp Asp Ala
    1625                1630                1635

Ile Thr  Ile Glu Thr Ile Glu  Asp Phe Asp Ile Tyr  Ser Glu Asp
    1640                1645                1650

Ile Lys  Gln Gly Pro Arg Ser  Phe Gln Gln Lys Thr  Arg His Tyr
    1655                1660                1665

Phe Ile  Ala Ala Val Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Thr
    1670                1675                1680

Ser His  Val Leu Arg Asn Arg  Tyr Gln Ser Asp Asn  Val Pro Gln
    1685                1690                1695

Phe Lys  Lys Val Val Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Ser
    1700                1705                1710
```

```
Gln Pro  Leu Tyr Arg Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu
    1715                1720                1725

Gly Pro  Tyr Ile Arg Ala Glu  Val Glu Asp Asn Ile  Met Val Thr
    1730                1735                1740

Phe Lys  Asn Gln Ala Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu
    1745                1750                1755

Ile Ser  Tyr Lys Glu Asp Gln  Arg Gly Glu Glu Pro  Arg Arg Asn
    1760                1765                1770

Phe Val  Lys Pro Asn Glu Thr  Lys Ile Tyr Phe Trp  Lys Val Gln
    1775                1780                1785

His His  Met Ala Pro Thr Glu  Asp Glu Phe Asp Cys  Lys Ala Trp
    1790                1795                1800

Ala Tyr  Phe Ser Asp Val Asp  Leu Glu Arg Asp Met  His Ser Gly
    1805                1810                1815

Leu Ile  Gly Pro Leu Leu Ile  Cys His Ala Asn Thr  Leu Asn Pro
    1820                1825                1830

Ala His  Gly Arg Gln Val Ser  Val Gln Glu Phe Ala  Leu Leu Phe
    1835                1840                1845

Thr Ile  Phe Asp Glu Thr Lys  Ser Trp Tyr Phe Thr  Glu Asn Val
    1850                1855                1860

Lys Arg  Asn Cys Lys Thr Pro  Cys Asn Phe Gln Met  Glu Asp Pro
    1865                1870                1875

Thr Leu  Lys Glu Asn Tyr Arg  Phe His Ala Ile Asn  Gly Tyr Val
    1880                1885                1890

Met Asp  Thr Leu Pro Gly Leu  Val Met Ala Gln Asp  Gln Arg Ile
    1895                1900                1905

Arg Trp  Tyr Leu Leu Ser Met  Gly Asn Asn Glu Asn  Ile Gln Ser
    1910                1915                1920

Ile His  Phe Ser Gly His Val  Phe Thr Val Arg Lys  Lys Glu Glu
    1925                1930                1935

Tyr Lys  Met Ala Val Tyr Asn  Leu Tyr Pro Gly Val  Phe Glu Thr
    1940                1945                1950

Leu Glu  Met Ile Pro Ser Arg  Ala Gly Ile Trp Arg  Val Glu Cys
    1955                1960                1965
```

```
Leu Ile  Gly Glu His Leu Gln  Ala Gly Met Ser Thr  Leu Phe Leu
    1970                1975                1980

Val Tyr  Ser Lys Gln Cys Gln  Ile Pro Leu Gly Met  Ala Ser Gly
    1985                1990                1995

Ser Ile  Arg Asp Phe Gln Ile  Thr Ala Ser Gly His  Tyr Gly Gln
    2000                2005                2010

Trp Ala  Pro Asn Leu Ala Arg  Leu His Tyr Ser Gly  Ser Ile Asn
    2015                2020                2025

Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp Ile Lys  Val Asp Leu
    2030                2035                2040

Leu Ala  Pro Met Ile Val His  Gly Ile Lys Thr Gln  Gly Ala Arg
    2045                2050                2055

Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln Phe Ile  Ile Met Tyr
    2060                2065                2070

Ser Leu  Asp Gly Lys Lys Trp  Leu Ser Tyr Gln Gly  Asn Ser Thr
    2075                2080                2085

Gly Thr  Leu Met Val Phe Phe  Gly Asn Val Asp Ser  Ser Gly Ile
    2090                2095                2100

Lys His  Asn Ser Phe Asn Pro  Pro Ile Ile Ala Arg  Tyr Ile Arg
    2105                2110                2115

Leu His  Pro Thr His Ser Ser  Ile Arg Ser Thr Leu  Arg Met Glu
    2120                2125                2130

Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser Ile Pro  Leu Gly Met
    2135                2140                2145

Glu Ser  Lys Val Ile Ser Asp  Thr Gln Ile Thr Ala  Ser Ser Tyr
    2150                2155                2160

Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro Ser Gln  Ala Arg Leu
    2165                2170                2175

His Leu  Gln Gly Arg Thr Asn  Ala Trp Arg Pro Gln  Val Asn Asp
    2180                2185                2190

Pro Lys  Gln Trp Leu Gln Val  Asp Leu Gln Lys Thr  Met Lys Val
    2195                2200                2205

Thr Gly  Ile Ile Thr Gln Gly  Val Lys Ser Leu Phe  Thr Ser Met
    2210                2215                2220
```

Phe Val Lys Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly His His
2225 2230 2235

Trp Thr Gln Ile Leu Tyr Asn Gly Lys Val Lys Val Phe Gln Gly
2240 2245 2250

Asn Gln Asp Ser Ser Thr Pro Met Met Asn Ser Leu Asp Pro Pro
2255 2260 2265

Leu Leu Thr Arg Tyr Leu Arg Ile His Pro Gln Ile Trp Glu His
2270 2275 2280

Gln Ile Ala Leu Arg Leu Glu Ile Leu Gly Cys Glu Ala Gln Gln
2285 2290 2295

Gln Tyr
2300

<210> 6
<211> 1457
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 6
Met Gln Ile Glu Leu Ser Thr Cys Phe Phe Leu Cys Leu Leu Arg Phe
1 5 10 15

Cys Phe Ser Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser
20 25 30

Trp Asp Tyr Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg
35 40 45

Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val
50 55 60

Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile
65 70 75 80

Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln
85 90 95

Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser
100 105 110

His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser
115 120 125

Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp
130 135 140

```
Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu
145                 150                 155                 160

Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser
                165                 170                 175

Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile
            180                 185                 190

Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr
        195                 200                 205

Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly
    210                 215                 220

Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp
225                 230                 235                 240

Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr
                245                 250                 255

Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val
            260                 265                 270

Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile
        275                 280                 285

Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser
    290                 295                 300

Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met
305                 310                 315                 320

Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His
                325                 330                 335

Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro
            340                 345                 350

Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp
        355                 360                 365

Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser
    370                 375                 380

Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr
385                 390                 395                 400

Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro
                405                 410                 415
```

```
Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn
            420                 425                 430

Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met
        435                 440                 445

Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu
    450                 455                 460

Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu
465                 470                 475                 480

Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro
                485                 490                 495

His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys
            500                 505                 510

Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe
        515                 520                 525

Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp
    530                 535                 540

Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg
545                 550                 555                 560

Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu
                565                 570                 575

Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val
            580                 585                 590

Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu
        595                 600                 605

Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp
    610                 615                 620

Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val
625                 630                 635                 640

Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp
                645                 650                 655

Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe
            660                 665                 670

Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr
        675                 680                 685
```

```
Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro
    690                 695                 700

Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly
705                 710                 715                 720

Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp
                725                 730                 735

Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys
            740                 745                 750

Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu
        755                 760                 765

Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln
    770                 775                 780

Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu
785                 790                 795                 800

Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe
                805                 810                 815

Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp
            820                 825                 830

Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln
        835                 840                 845

Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr
    850                 855                 860

Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His
865                 870                 875                 880

Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile
                885                 890                 895

Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser
            900                 905                 910

Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg
        915                 920                 925

Lys Asn Phe Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val
    930                 935                 940

Gln His His Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp
945                 950                 955                 960
```

```
Ala Tyr Phe Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu
                965                 970                 975

Ile Gly Pro Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His
            980                 985                 990

Gly Arg Gln Val Thr Val Gln Glu  Phe Ala Leu Phe Phe  Thr Ile Phe
        995                 1000                1005

Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu Asn Met  Glu Arg Asn
    1010                1015                1020

Cys Arg  Ala Pro Cys Asn Ile  Gln Met Glu Asp Pro  Thr Phe Lys
    1025                1030                1035

Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly Tyr Ile  Met Asp Thr
    1040                1045                1050

Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln Arg Ile  Arg Trp Tyr
    1055                1060                1065

Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile His Ser  Ile His Phe
    1070                1075                1080

Ser Gly  His Val Phe Thr Val  Arg Lys Lys Glu Glu  Tyr Lys Met
    1085                1090                1095

Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe Glu Thr  Val Glu Met
    1100                1105                1110

Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val Glu Cys  Leu Ile Gly
    1115                1120                1125

Glu His  Leu His Ala Gly Met  Ser Thr Leu Phe Leu  Val Tyr Ser
    1130                1135                1140

Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala Ser Gly  His Ile Arg
    1145                1150                1155

Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr Gly Gln  Trp Ala Pro
    1160                1165                1170

Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser Ile Asn  Ala Trp Ser
    1175                1180                1185

Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val Asp Leu  Leu Ala Pro
    1190                1195                1200

Met Ile  Ile His Gly Ile Lys  Thr Gln Gly Ala Arg  Gln Lys Phe
    1205                1210                1215
```

```
Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile Met Tyr  Ser Leu Asp
    1220                 1225                 1230

Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn Ser Thr  Gly Thr Leu
    1235                 1240                 1245

Met Val  Phe Phe Gly Asn Val  Asp Ser Ser Gly Ile  Lys His Asn
    1250                 1255                 1260

Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr Ile Arg  Leu His Pro
    1265                 1270                 1275

Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg Met Glu  Leu Met Gly
    1280                 1285                 1290

Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu Gly Met  Glu Ser Lys
    1295                 1300                 1305

Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser Ser Tyr  Phe Thr Asn
    1310                 1315                 1320

Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala Arg Leu  His Leu Gln
    1325                 1330                 1335

Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val Asn Asn  Pro Lys Glu
    1340                 1345                 1350

Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met Lys Val  Thr Gly Val
    1355                 1360                 1365

Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr Ser Met  Tyr Val Lys
    1370                 1375                 1380

Glu Phe  Leu Ile Ser Ser Ser  Gln Asp Gly His Gln  Trp Thr Leu
    1385                 1390                 1395

Phe Phe  Gln Asn Gly Lys Val  Lys Val Phe Gln Gly  Asn Gln Asp
    1400                 1405                 1410

Ser Phe  Thr Pro Val Val Asn  Ser Leu Asp Pro Pro  Leu Leu Thr
    1415                 1420                 1425

Arg Tyr  Leu Arg Ile His Pro  Gln Ser Trp Val His  Gln Ile Ala
    1430                 1435                 1440

Leu Arg  Met Glu Val Leu Gly  Cys Glu Ala Gln Asp  Leu Tyr
    1445                 1450                 1455
```

<210> 7
<211> 1438

<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 7

```
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1               5                  10                  15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240
```

```
Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510
```

```
Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780
```

```
Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1025                1030                1035

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1040                1045                1050
```

```
Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1055                1060                1065

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1070                1075                1080

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1085                1090                1095

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1100                1105                1110

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1115                1120                1125

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1130                1135                1140

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1145                1150                1155

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1160                1165                1170

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1175                1180                1185

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1190                1195                1200

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1205                1210                1215

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1220                1225                1230

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1235                1240                1245

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1250                1255                1260

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1265                1270                1275

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1280                1285                1290

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1295                1300                1305
```

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1310 1315 1320

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1325 1330 1335

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1340 1345 1350

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1355 1360 1365

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1370 1375 1380

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1385 1390 1395

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1400 1405 1410

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1415 1420 1425

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr
1430 1435

<210> 8
<211> 1438
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 8
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

```
Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365
```

```
Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640
```

```
Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Gly Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910
```

```
Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1025                1030                1035

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1040                1045                1050

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1055                1060                1065

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1070                1075                1080

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1085                1090                1095

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1100                1105                1110

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1115                1120                1125

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1130                1135                1140

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1145                1150                1155

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1160                1165                1170
```

```
Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1175                1180                1185

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1190                1195                1200

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1205                1210                1215

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1220                1225                1230

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1235                1240                1245

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1250                1255                1260

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1265                1270                1275

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1280                1285                1290

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1295                1300                1305

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1310                1315                1320

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1325                1330                1335

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1340                1345                1350

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1355                1360                1365

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1370                1375                1380

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1385                1390                1395

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1400                1405                1410

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1415                1420                1425
```

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr
    1430   1435

<210> 9
<211> 1391
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
      polypeptide

<400> 9
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1   5   10   15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20   25   30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35   40   45

Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro
    50   55   60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65   70   75   80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85   90   95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100   105   110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115   120   125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130   135   140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145   150   155   160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165   170   175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180   185   190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195   200   205

```
His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480
```

```
Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Gln Ser Pro Arg Ser Phe Gln Lys
            740                 745                 750
```

```
Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr
        755                 760                 765

Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly
    770                 775                 780

Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly
785                 790                 795                 800

Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly
                805                 810                 815

Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val
            820                 825                 830

Thr Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu
        835                 840                 845

Ile Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn
    850                 855                 860

Phe Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His
865                 870                 875                 880

His Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr
                885                 890                 895

Phe Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly
            900                 905                 910

Pro Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg
        915                 920                 925

Gln Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu
    930                 935                 940

Thr Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala
945                 950                 955                 960

Pro Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
                965                 970                 975

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu Val
            980                 985                 990

Met Ala Gln Asp Gln Arg Ile Arg  Trp Tyr Leu Leu Ser  Met Gly Ser
        995                 1000                1005

Asn Glu  Asn Ile His Ser Ile  His Phe Ser Gly His  Val Phe Thr
    1010                1015                1020
```

```
Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala Leu Tyr  Asn Leu Tyr
    1025                1030                1035

Pro Gly  Val Phe Glu Thr Val  Glu Met Leu Pro Ser  Lys Ala Gly
    1040                1045                1050

Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu His Leu  His Ala Gly
    1055                1060                1065

Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn Lys Cys  Gln Thr Pro
    1070                1075                1080

Leu Gly  Met Ala Ser Gly His  Ile Arg Asp Phe Gln  Ile Thr Ala
    1085                1090                1095

Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys Leu Ala  Arg Leu His
    1100                1105                1110

Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr Lys Glu  Pro Phe Ser
    1115                1120                1125

Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met Ile Ile  His Gly Ile
    1130                1135                1140

Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser Ser Leu  Tyr Ile Ser
    1145                1150                1155

Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly Lys Lys  Trp Gln Thr
    1160                1165                1170

Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met Val Phe  Phe Gly Asn
    1175                1180                1185

Val Asp  Ser Ser Gly Ile Lys  His Asn Ile Phe Asn  Pro Pro Ile
    1190                1195                1200

Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr His Tyr  Ser Ile Arg
    1205                1210                1215

Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys Asp Leu  Asn Ser Cys
    1220                1225                1230

Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala Ile Ser  Asp Ala Gln
    1235                1240                1245

Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met Phe Ala  Thr Trp Ser
    1250                1255                1260

Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly Arg Ser  Asn Ala Trp
    1265                1270                1275
```

Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe
1280 1285 1290

Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys
1295 1300 1305

Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser
1310 1315 1320

Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys
1325 1330 1335

Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val
1340 1345 1350

Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His
1355 1360 1365

Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu
1370 1375 1380

Gly Cys Glu Ala Gln Asp Leu Tyr
1385 1390

<210> 10
<211> 1360
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 10
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

```
Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380
```

```
Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655
```

```
Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Gln Ser Pro Arg Ser Phe Gln
705                 710                 715                 720

Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp
                725                 730                 735

Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser
            740                 745                 750

Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp
        755                 760                 765

Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu
    770                 775                 780

Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met
785                 790                 795                 800

Val Thr Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser
                805                 810                 815

Leu Ile Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys
            820                 825                 830

Asn Phe Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln
        835                 840                 845

His His Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala
    850                 855                 860

Tyr Phe Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile
865                 870                 875                 880

Gly Pro Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly
                885                 890                 895

Arg Gln Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp
            900                 905                 910

Glu Thr Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg
        915                 920                 925
```

```
Ala Pro Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr
    930                 935                 940

Arg Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
945                 950                 955                 960

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met Gly
                965                 970                 975

Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val Phe Thr
            980                 985                 990

Val Arg Lys Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
        995                 1000                1005

Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    1010                1015                1020

Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    1025                1030                1035

Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    1040                1045                1050

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
    1055                1060                1065

Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
    1070                1075                1080

Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
    1085                1090                1095

Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
    1100                1105                1110

Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    1115                1120                1125

Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
    1130                1135                1140

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    1145                1150                1155

Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
    1160                1165                1170

Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1175                1180                1185
```

```
Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
    1190                1195                1200

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    1205                1210                1215

Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
    1220                1225                1230

Ser Lys  Ala Arg Leu His Leu  Gln Gly Arg Ser Asn  Ala Trp Arg
    1235                1240                1245

Pro Gln  Val Asn Asn Pro Lys  Glu Trp Leu Gln Val  Asp Phe Gln
    1250                1255                1260

Lys Thr  Met Lys Val Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser
    1265                1270                1275

Leu Leu  Thr Ser Met Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser
    1280                1285                1290

Gln Asp  Gly His Gln Trp Thr  Leu Phe Phe Gln Asn  Gly Lys Val
    1295                1300                1305

Lys Val  Phe Gln Gly Asn Gln  Asp Ser Phe Thr Pro  Val Val Asn
    1310                1315                1320

Ser Leu  Asp Pro Pro Leu Leu  Thr Arg Tyr Leu Arg  Ile His Pro
    1325                1330                1335

Gln Ser  Trp Val His Gln Ile  Ala Leu Arg Met Glu  Val Leu Gly
    1340                1345                1350

Cys Glu  Ala Gln Asp Leu Tyr
    1355                1360
```

<210> 11
<211> 1436
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 11

```
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1               5                  10                  15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30
```

```
Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300
```

```
Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575
```

```
Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Ser Arg His Pro Ser Thr Arg
            740                 745                 750

Gln Lys Gln Phe Asn Ala Thr Thr Ile Pro Glu Asn Asp Ile Glu Lys
        755                 760                 765

Thr Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr
    770                 775                 780

Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg
785                 790                 795                 800

His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser
                805                 810                 815

Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro
            820                 825                 830

Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr
        835                 840                 845
```

```
Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly
    850                 855                 860

Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg
865                 870                 875                 880

Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr
                885                 890                 895

Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys
            900                 905                 910

Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala
        915                 920                 925

Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp
    930                 935                 940

Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu
945                 950                 955                 960

Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
                965                 970                 975

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys Ser
            980                 985                 990

Trp Tyr Phe Thr Glu Asn Met Glu  Arg Asn Cys Arg Ala  Pro Cys Asn
        995                 1000                1005

Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu Asn Tyr  Arg Phe His
    1010                1015                1020

Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu Pro Gly  Leu Val Met
    1025                1030                1035

Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu Leu Ser  Met Gly Ser
    1040                1045                1050

Asn Glu  Asn Ile His Ser Ile  His Phe Ser Gly His  Val Phe Thr
    1055                1060                1065

Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala Leu Tyr  Asn Leu Tyr
    1070                1075                1080

Pro Gly  Val Phe Glu Thr Val  Glu Met Leu Pro Ser  Lys Ala Gly
    1085                1090                1095

Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu His Leu  His Ala Gly
    1100                1105                1110
```

```
Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn Lys Cys  Gln Thr Pro
    1115                1120                1125

Leu Gly  Met Ala Ser Gly His  Ile Arg Asp Phe Gln  Ile Thr Ala
    1130                1135                1140

Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys Leu Ala  Arg Leu His
    1145                1150                1155

Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr Lys Glu  Pro Phe Ser
    1160                1165                1170

Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met Ile Ile  His Gly Ile
    1175                1180                1185

Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser Ser Leu  Tyr Ile Ser
    1190                1195                1200

Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly Lys Lys  Trp Gln Thr
    1205                1210                1215

Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met Val Phe  Phe Gly Asn
    1220                1225                1230

Val Asp  Ser Ser Gly Ile Lys  His Asn Ile Phe Asn  Pro Pro Ile
    1235                1240                1245

Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr His Tyr  Ser Ile Arg
    1250                1255                1260

Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys Asp Leu  Asn Ser Cys
    1265                1270                1275

Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala Ile Ser  Asp Ala Gln
    1280                1285                1290

Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met Phe Ala  Thr Trp Ser
    1295                1300                1305

Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly Arg Ser  Asn Ala Trp
    1310                1315                1320

Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp Leu Gln  Val Asp Phe
    1325                1330                1335

Gln Lys  Thr Met Lys Val Thr  Gly Val Thr Thr Gln  Gly Val Lys
    1340                1345                1350

Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu Phe Leu  Ile Ser Ser
    1355                1360                1365
```

Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys
1370 1375 1380

Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val
1385 1390 1395

Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His
1400 1405 1410

Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu
1415 1420 1425

Gly Cys Glu Ala Gln Asp Leu Tyr
1430 1435

<210> 12
<211> 1390
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 12
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

```
His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430
```

```
Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700
```

```
Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Ser Pro Arg Ser Phe Gln Lys Lys
            740                 745                 750

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
        755                 760                 765

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
    770                 775                 780

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
785                 790                 795                 800

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
                805                 810                 815

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
            820                 825                 830

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
        835                 840                 845

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
    850                 855                 860

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
865                 870                 875                 880

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
                885                 890                 895

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
            900                 905                 910

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
        915                 920                 925

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
    930                 935                 940

Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
945                 950                 955                 960

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg Phe
                965                 970                 975
```

```
His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu Val Met
           980                 985                 990

Ala Gln Asp Gln Arg Ile Arg Trp  Tyr Leu Leu Ser Met  Gly Ser Asn
       995                 1000                1005

Glu Asn  Ile His Ser Ile His  Phe Ser Gly His Val  Phe Thr Val
    1010                1015                1020

Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
    1025                1030                1035

Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    1040                1045                1050

Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    1055                1060                1065

Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    1070                1075                1080

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
    1085                1090                1095

Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
    1100                1105                1110

Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
    1115                1120                1125

Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
    1130                1135                1140

Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    1145                1150                1155

Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
    1160                1165                1170

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    1175                1180                1185

Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
    1190                1195                1200

Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1205                1210                1215

Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
    1220                1225                1230
```

Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp Ala Gln Ile
1235 1240 1245

Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp Ser Pro
1250 1255 1260

Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg
1265 1270 1275

Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln
1280 1285 1290

Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser
1295 1300 1305

Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser
1310 1315 1320

Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val
1325 1330 1335

Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn
1340 1345 1350

Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro
1355 1360 1365

Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly
1370 1375 1380

Cys Glu Ala Gln Asp Leu Tyr
1385 1390

<210> 13
<211> 1457
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 13
Met Gln Ile Glu Leu Ser Thr Cys Phe Phe Leu Cys Leu Leu Arg Phe
1 5 10 15

Cys Phe Ser Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser
20 25 30

Trp Asp Tyr Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg
35 40 45

```
Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val
    50                  55                  60

Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile
65                  70                  75                  80

Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln
                85                  90                  95

Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser
            100                 105                 110

His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser
        115                 120                 125

Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp
    130                 135                 140

Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu
145                 150                 155                 160

Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser
                165                 170                 175

Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile
            180                 185                 190

Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr
        195                 200                 205

Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly
    210                 215                 220

Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp
225                 230                 235                 240

Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr
                245                 250                 255

Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val
            260                 265                 270

Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile
        275                 280                 285

Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser
    290                 295                 300

Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met
305                 310                 315                 320
```

```
Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His
                325                 330                 335

Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro
            340                 345                 350

Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp
        355                 360                 365

Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser
    370                 375                 380

Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr
385                 390                 395                 400

Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro
                405                 410                 415

Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn
            420                 425                 430

Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met
        435                 440                 445

Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu
    450                 455                 460

Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu
465                 470                 475                 480

Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro
                485                 490                 495

His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys
            500                 505                 510

Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe
        515                 520                 525

Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp
    530                 535                 540

Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg
545                 550                 555                 560

Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu
                565                 570                 575

Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val
            580                 585                 590
```

```
Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu
        595                 600                 605

Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp
    610                 615                 620

Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val
625                 630                 635                 640

Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp
                645                 650                 655

Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe
            660                 665                 670

Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr
        675                 680                 685

Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro
    690                 695                 700

Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly
705                 710                 715                 720

Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp
                725                 730                 735

Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys
            740                 745                 750

Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu
        755                 760                 765

Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln
    770                 775                 780

Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu
785                 790                 795                 800

Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe
                805                 810                 815

Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp
            820                 825                 830

Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln
        835                 840                 845

Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr
    850                 855                 860
```

```
Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His
865                 870                 875                 880

Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile
                885                 890                 895

Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser
            900                 905                 910

Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg
        915                 920                 925

Lys Asn Phe Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val
    930                 935                 940

Gln His His Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp
945                 950                 955                 960

Ala Tyr Phe Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu
                965                 970                 975

Ile Gly Pro Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His
            980                 985                 990

Gly Arg Gln Val Thr Val Gln Glu  Phe Ala Leu Phe Phe  Thr Ile Phe
        995                 1000                1005

Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu Asn Met  Glu Arg Asn
    1010                1015                1020

Cys Arg  Ala Pro Cys Asn Ile  Gln Met Glu Asp Pro  Thr Phe Lys
    1025                1030                1035

Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly Tyr Ile  Met Asp Thr
    1040                1045                1050

Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln Arg Ile  Arg Trp Tyr
    1055                1060                1065

Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile His Ser  Ile His Phe
    1070                1075                1080

Ser Gly  His Val Phe Thr Val  Arg Lys Lys Glu Glu  Tyr Lys Met
    1085                1090                1095

Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe Glu Thr  Val Glu Met
    1100                1105                1110

Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val Glu Cys  Leu Ile Gly
    1115                1120                1125
```

```
Glu His  Leu His Ala Gly Met  Ser Thr Leu Phe Leu  Val Tyr Ser
    1130                1135                1140

Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala Ser Gly  His Ile Arg
    1145                1150                1155

Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr Gly Gln  Trp Ala Pro
    1160                1165                1170

Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser Ile Asn  Ala Trp Ser
    1175                1180                1185

Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val Asp Leu  Leu Ala Pro
    1190                1195                1200

Met Ile  Ile His Gly Ile Lys  Thr Gln Gly Ala Arg  Gln Lys Phe
    1205                1210                1215

Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile Met Tyr  Ser Leu Asp
    1220                1225                1230

Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn Ser Thr  Gly Thr Leu
    1235                1240                1245

Met Val  Phe Phe Gly Asn Val  Asp Ser Ser Gly Ile  Lys His Asn
    1250                1255                1260

Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr Ile Arg  Leu His Pro
    1265                1270                1275

Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg Met Glu  Leu Met Gly
    1280                1285                1290

Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu Gly Met  Glu Ser Lys
    1295                1300                1305

Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser Ser Tyr  Phe Thr Asn
    1310                1315                1320

Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala Arg Leu  His Leu Gln
    1325                1330                1335

Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val Asn Asn  Pro Lys Glu
    1340                1345                1350

Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met Lys Val  Thr Gly Val
    1355                1360                1365

Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr Ser Met  Tyr Val Lys
    1370                1375                1380
```

Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu
1385 1390 1395

Phe Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp
1400 1405 1410

Ser Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr
1415 1420 1425

Arg Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile Ala
1430 1435 1440

Leu Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr
1445 1450 1455

<210> 14
<211> 1438
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 14
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

```
His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430
```

```
Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700
```

```
Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975
```

```
Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1025                1030                1035

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1040                1045                1050

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1055                1060                1065

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1070                1075                1080

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1085                1090                1095

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1100                1105                1110

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1115                1120                1125

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1130                1135                1140

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1145                1150                1155

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1160                1165                1170

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1175                1180                1185

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1190                1195                1200

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1205                1210                1215

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1220                1225                1230
```

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1235 1240 1245

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1250 1255 1260

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1265 1270 1275

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1280 1285 1290

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1295 1300 1305

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1310 1315 1320

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1325 1330 1335

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1340 1345 1350

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1355 1360 1365

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1370 1375 1380

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1385 1390 1395

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1400 1405 1410

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1415 1420 1425

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr
1430 1435

<210> 15
<211> 1438
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 15

```
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1               5                  10                  15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270
```

```
Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540
```

```
Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815
```

```
Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1025                1030                1035

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1040                1045                1050

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1055                1060                1065

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1070                1075                1080
```

```
Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1085                1090                1095

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1100                1105                1110

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1115                1120                1125

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1130                1135                1140

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1145                1150                1155

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1160                1165                1170

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1175                1180                1185

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1190                1195                1200

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1205                1210                1215

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1220                1225                1230

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1235                1240                1245

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1250                1255                1260

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1265                1270                1275

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1280                1285                1290

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1295                1300                1305

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1310                1315                1320

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1325                1330                1335
```

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1340 1345 1350

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1355 1360 1365

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1370 1375 1380

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1385 1390 1395

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1400 1405 1410

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1415 1420 1425

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr
1430 1435

<210> 16
<211> 1441
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 16
Ala Thr Arg Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser
1 5 10 15

Trp Asp Tyr Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg
20 25 30

Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val
35 40 45

Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile
50 55 60

Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln
65 70 75 80

Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser
85 90 95

His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser
100 105 110

Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp
115 120 125

```
Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu
    130                 135                 140

Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser
145                 150                 155                 160

Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile
                165                 170                 175

Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr
            180                 185                 190

Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly
        195                 200                 205

Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp
    210                 215                 220

Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr
225                 230                 235                 240

Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val
                245                 250                 255

Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile
            260                 265                 270

Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser
        275                 280                 285

Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met
    290                 295                 300

Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His
305                 310                 315                 320

Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro
                325                 330                 335

Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp
            340                 345                 350

Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser
        355                 360                 365

Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr
    370                 375                 380

Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro
385                 390                 395                 400
```

```
Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn
                405                 410                 415

Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met
            420                 425                 430

Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu
        435                 440                 445

Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu
    450                 455                 460

Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro
465                 470                 475                 480

His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys
                485                 490                 495

Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe
            500                 505                 510

Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp
        515                 520                 525

Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg
    530                 535                 540

Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu
545                 550                 555                 560

Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val
                565                 570                 575

Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu
            580                 585                 590

Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp
        595                 600                 605

Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val
    610                 615                 620

Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp
625                 630                 635                 640

Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe
                645                 650                 655

Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr
            660                 665                 670
```

```
Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro
        675                 680                 685

Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly
    690                 695                 700

Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp
705                 710                 715                 720

Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys
                725                 730                 735

Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu
            740                 745                 750

Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln
        755                 760                 765

Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu
    770                 775                 780

Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe
785                 790                 795                 800

Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp
                805                 810                 815

Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln
            820                 825                 830

Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr
        835                 840                 845

Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His
    850                 855                 860

Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile
865                 870                 875                 880

Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser
                885                 890                 895

Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg
            900                 905                 910

Lys Asn Phe Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val
        915                 920                 925

Gln His His Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp
    930                 935                 940
```

```
Ala Tyr Phe Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu
945                 950                 955                 960

Ile Gly Pro Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His
                965                 970                 975

Gly Arg Gln Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe
            980                 985                 990

Asp Glu Thr Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
        995                 1000                1005

Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1010                1015                1020

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1025                1030                1035

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1040                1045                1050

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1055                1060                1065

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1070                1075                1080

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1085                1090                1095

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1100                1105                1110

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1115                1120                1125

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1130                1135                1140

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1145                1150                1155

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1160                1165                1170

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1175                1180                1185

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1190                1195                1200
```

```
Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1205                 1210                 1215

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1220                 1225                 1230

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1235                 1240                 1245

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1250                 1255                 1260

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1265                 1270                 1275

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1280                 1285                 1290

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1295                 1300                 1305

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1310                 1315                 1320

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1325                 1330                 1335

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1340                 1345                 1350

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1355                 1360                 1365

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1370                 1375                 1380

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1385                 1390                 1395

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    1400                 1405                 1410

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    1415                 1420                 1425

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr
    1430                 1435                 1440
```

<210> 17
<211> 1438

<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 17

```
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1               5                   10                  15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240
```

```
Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
               245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
               325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
               405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
               485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510
```

```
Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780
```

```
Ile Phe Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1025                1030                1035

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1040                1045                1050
```

```
Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1055                1060                1065

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1070                1075                1080

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1085                1090                1095

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1100                1105                1110

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1115                1120                1125

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1130                1135                1140

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1145                1150                1155

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1160                1165                1170

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1175                1180                1185

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1190                1195                1200

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1205                1210                1215

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1220                1225                1230

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1235                1240                1245

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1250                1255                1260

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1265                1270                1275

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1280                1285                1290

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1295                1300                1305
```

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1310 1315 1320

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1325 1330 1335

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1340 1345 1350

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1355 1360 1365

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1370 1375 1380

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1385 1390 1395

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1400 1405 1410

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1415 1420 1425

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr
1430 1435

<210> 18
<211> 1438
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 18
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

```
Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365
```

```
Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640
```

```
Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Phe Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910
```

```
Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1025                1030                1035

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1040                1045                1050

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1055                1060                1065

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1070                1075                1080

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1085                1090                1095

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1100                1105                1110

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1115                1120                1125

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1130                1135                1140

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1145                1150                1155

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1160                1165                1170
```

```
Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1175                1180                1185

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1190                1195                1200

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1205                1210                1215

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1220                1225                1230

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1235                1240                1245

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1250                1255                1260

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1265                1270                1275

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1280                1285                1290

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1295                1300                1305

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1310                1315                1320

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1325                1330                1335

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1340                1345                1350

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1355                1360                1365

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1370                1375                1380

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1385                1390                1395

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1400                1405                1410

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1415                1420                1425
```

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr
    1430   1435

<210> 19
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
      peptide

<400> 19
Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser
1   5   10

<210> 20
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
      peptide

<400> 20
Gly Ser Glu Gly Ser Ser Gly Pro Gly Glu Ser Ser
1   5   10

<210> 21
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
      peptide

<400> 21
Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro
1   5   10

<210> 22
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
      peptide

<400> 22
Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser
1   5   10

<210> 23
<211> 12
<212> PRT
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic peptide

<400> 23
Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
1 5 10

<210> 24
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 24
Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
1 5 10

<210> 25
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 25
Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
1 5 10

<210> 26
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 26
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
1 5 10

<210> 27
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 27
Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
1 5 10

<210> 28
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 28
Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
1 5 10

<210> 29
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 29
Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro
1 5 10

<210> 30
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 30
Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro
1 5 10

<210> 31
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 31
Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
1 5 10

<210> 32
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 32
Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
1 5 10

<210> 33
<211> 12

<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 33
Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
1 5 10

<210> 34
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 34
Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
1 5 10

<210> 35
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 35
Gly Glu Pro Ala Gly Ser Pro Thr Ser Thr Ser Glu
1 5 10

<210> 36
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 36
Gly Thr Gly Glu Pro Ser Ser Thr Pro Ala Ser Glu
1 5 10

<210> 37
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 37
Gly Ser Gly Pro Ser Thr Glu Ser Ala Pro Thr Glu
1 5 10

<210> 38
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 38
Gly Ser Glu Thr Pro Ser Gly Pro Ser Glu Thr Ala
1 5 10

<210> 39
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 39
Gly Pro Ser Glu Thr Ser Thr Ser Glu Pro Gly Ala
1 5 10

<210> 40
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 40
Gly Ser Pro Ser Glu Pro Thr Glu Gly Thr Ser Ala
1 5 10

<210> 41
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 41
Gly Ser Gly Ala Ser Glu Pro Thr Ser Thr Glu Pro
1 5 10

<210> 42
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 42
Gly Ser Glu Pro Ala Thr Ser Gly Thr Glu Pro Ser
1 5 10

<210> 43
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 43
Gly Thr Ser Glu Pro Ser Thr Ser Glu Pro Gly Ala
1 5 10

<210> 44
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 44
Gly Thr Ser Thr Glu Pro Ser Glu Pro Gly Ser Ala
1 5 10

<210> 45
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 45
Gly Ser Thr Ala Gly Ser Glu Thr Ser Thr Glu Ala
1 5 10

<210> 46
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 46
Gly Ser Glu Thr Ala Thr Ser Gly Ser Glu Thr Ala
1 5 10

<210> 47
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 47

Gly Thr Ser Glu Ser Ala Thr Ser Glu Ser Gly Ala
1 5 10

<210> 48
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 48
Gly Thr Ser Thr Glu Ala Ser Glu Gly Ser Ala Ser
1 5 10

<210> 49
<211> 42
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 49
Gly Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
1 5 10 15

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
20 25 30

Thr Ser Gly Ser Glu Thr Pro Ala Ser Ser
35 40

<210> 50
<211> 42
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 50
Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro
1 5 10 15

Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
20 25 30

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
35 40

<210> 51
<211> 42
<212> PRT
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 51
Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro
1 5 10 15

Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
20 25 30

Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
35 40

<210> 52
<211> 42
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 52
Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
1 5 10 15

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
20 25 30

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro
35 40

<210> 53
<211> 42
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 53
Gly Ala Pro Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
1 5 10 15

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
20 25 30

Ala Thr Gly Ser Pro Gly Pro Ser Gly Pro
35 40

<210> 54
<211> 42
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 54
Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser
1 5 10 15

Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala
20 25 30

Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
35 40

<210> 55
<211> 42
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 55
Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr
1 5 10 15

Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
20 25 30

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
35 40

<210> 56
<211> 42
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 56
Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
1 5 10 15

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
20 25 30

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
35 40

<210> 57
<211> 48
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 57

Met Ala Glu Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Pro
1 5 10 15

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
20 25 30

Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
35 40 45

<210> 58
<211> 48
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 58
Met Ala Glu Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ala Ser
1 5 10 15

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly
20 25 30

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
35 40 45

<210> 59
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 59
Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
20 25 30

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
35 40 45

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
50 55 60

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
65 70 75 80

Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
85 90 95

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
100 105 110

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
115 120 125

Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
130 135 140

<210> 60
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 60
Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
1 5 10 15

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
20 25 30

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
35 40 45

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
50 55 60

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
65 70 75 80

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
85 90 95

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly
100 105 110

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
115 120 125

Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
130 135 140

<210> 61
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 61

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
1 5 10 15

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
20 25 30

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
35 40 45

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
50 55 60

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
65 70 75 80

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
85 90 95

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
100 105 110

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
115 120 125

Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
130 135 140

<210> 62
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 62
Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
1 5 10 15

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
20 25 30

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
35 40 45

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
50 55 60

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
65 70 75 80

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
85 90 95

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
100 105 110

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
115 120 125

Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
130 135 140

<210> 63
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 63
Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
1 5 10 15

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
20 25 30

Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
35 40 45

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
50 55 60

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
65 70 75 80

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
85 90 95

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
100 105 110

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
115 120 125

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
130 135 140

<210> 64
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 64
Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
1 5 10 15

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
20 25 30

Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
35 40 45

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
50 55 60

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
65 70 75 80

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
85 90 95

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
100 105 110

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
115 120 125

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
130 135 140

<210> 65
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 65
Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1 5 10 15

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
20 25 30

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
35 40 45

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
50 55 60

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
65 70 75 80

Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
85 90 95

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
100 105 110

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
115 120 125

Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
130 135 140

<210> 66
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 66
Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1 5 10 15

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
20 25 30

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
35 40 45

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
50 55 60

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
65 70 75 80

Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
85 90 95

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
100 105 110

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
115 120 125

Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
130 135 140

<210> 67
<211> 144
<212> PRT
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 67
Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1 5 10 15

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
20 25 30

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
35 40 45

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
50 55 60

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
65 70 75 80

Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
85 90 95

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
100 105 110

Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
115 120 125

Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
130 135 140

<210> 68
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 68
Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
1 5 10 15

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
20 25 30

Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
35 40 45

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala
50 55 60

Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
65 70 75 80

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
85 90 95

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
100 105 110

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
115 120 125

Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
130 135 140

<210> 69
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 69
Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Thr Ser Pro
1 5 10 15

Ser Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser
20 25 30

Ser Thr Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro
35 40 45

Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser
50 55 60

Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro Ser Gly Glu Ser
65 70 75 80

Ser Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
85 90 95

Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro
100 105 110

Ser Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser
115 120 125

Ser Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro
130 135 140

<210> 70
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 70
Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
1 5 10 15

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
20 25 30

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
35 40 45

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
50 55 60

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
65 70 75 80

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
85 90 95

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
100 105 110

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
115 120 125

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
130 135 140

<210> 71
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 71
Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser
1 5 10 15

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr
20 25 30

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
35 40 45

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
50 55 60

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
65 70 75 80

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
85 90 95

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser
100 105 110

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
115 120 125

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
130 135 140

<210> 72
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 72
Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
1 5 10 15

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
20 25 30

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
35 40 45

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
50 55 60

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
65 70 75 80

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
85 90 95

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
100 105 110

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
115 120 125

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
130 135 140

<210> 73
<211> 144

<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 73
Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
1 5 10 15

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
20 25 30

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
35 40 45

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
50 55 60

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
65 70 75 80

Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
85 90 95

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro
100 105 110

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
115 120 125

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
130 135 140

<210> 74
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 74
Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
1 5 10 15

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
20 25 30

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
35 40 45

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
50 55 60

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
65 70 75 80

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
85 90 95

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
100 105 110

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
115 120 125

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
130 135 140

<210> 75
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 75
Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
1 5 10 15

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
20 25 30

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
35 40 45

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
50 55 60

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
65 70 75 80

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
85 90 95

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
100 105 110

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
115 120 125

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
130 135 140

<210> 76
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 76
Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
1 5 10 15

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
20 25 30

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
35 40 45

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
50 55 60

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
65 70 75 80

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
85 90 95

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
100 105 110

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
115 120 125

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
130 135 140

<210> 77
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 77
Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
1 5 10 15

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
20 25 30

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
35 40 45

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
50 55 60

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
65 70 75 80

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
85 90 95

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
100 105 110

Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala
115 120 125

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
130 135 140

<210> 78
<211> 288
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 78
Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
1 5 10 15

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
20 25 30

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
35 40 45

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
50 55 60

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
65 70 75 80

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
85 90 95

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
100 105 110

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
115 120 125

Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
130 135 140

```
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
145                 150                 155                 160

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
                165                 170                 175

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
            180                 185                 190

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
        195                 200                 205

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
    210                 215                 220

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
225                 230                 235                 240

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
                245                 250                 255

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
            260                 265                 270

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
        275                 280                 285


<210> 79
<211> 288
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
      polypeptide

<400> 79
Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
1               5                   10                  15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
            20                  25                  30

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        35                  40                  45

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
    50                  55                  60

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
65                  70                  75                  80
```

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
85 90 95

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
100 105 110

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
115 120 125

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
130 135 140

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
145 150 155 160

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
165 170 175

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
180 185 190

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
195 200 205

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
210 215 220

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
225 230 235 240

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala
245 250 255

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
260 265 270

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
275 280 285

<210> 80
<211> 288
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 80
Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
1 5 10 15

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
20 25 30

```
Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
        35                  40                  45

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser
    50                  55                  60

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
65                  70                  75                  80

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
                85                  90                  95

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
            100                 105                 110

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
        115                 120                 125

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
    130                 135                 140

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser
145                 150                 155                 160

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
                165                 170                 175

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
            180                 185                 190

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
        195                 200                 205

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly
    210                 215                 220

Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
225                 230                 235                 240

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
                245                 250                 255

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr
            260                 265                 270

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
        275                 280                 285
```

<210> 81
<211> 288

<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 81

```
Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
1               5                   10                  15

Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala
            20                  25                  30

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
        35                  40                  45

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
    50                  55                  60

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
65                  70                  75                  80

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
                85                  90                  95

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
            100                 105                 110

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
        115                 120                 125

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
    130                 135                 140

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro
145                 150                 155                 160

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
                165                 170                 175

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
            180                 185                 190

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
        195                 200                 205

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala
    210                 215                 220

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
225                 230                 235                 240
```

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
245 250 255

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
260 265 270

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
275 280 285

<210> 82
<211> 504
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<220>
<221> MOD_RES
<222> (63)..(63)
<223> Any amino acid

<220>
<221> MOD_RES
<222> (207)..(207)
<223> Any amino acid

<400> 82
Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
1 5 10 15

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
20 25 30

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
35 40 45

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Xaa Pro
50 55 60

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
65 70 75 80

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
85 90 95

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
100 105 110

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
115 120 125

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
130 135 140

```
Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
145                 150                 155                 160

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
                165                 170                 175

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
            180                 185                 190

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Xaa Pro
        195                 200                 205

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
    210                 215                 220

Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
225                 230                 235                 240

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro
                245                 250                 255

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
            260                 265                 270

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
        275                 280                 285

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
    290                 295                 300

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
305                 310                 315                 320

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                325                 330                 335

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
            340                 345                 350

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
        355                 360                 365

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    370                 375                 380

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
385                 390                 395                 400

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
                405                 410                 415
```

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
420 425 430

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
435 440 445

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
450 455 460

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
465 470 475 480

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
485 490 495

Gly Thr Ser Ser Thr Gly Ser Pro
500

<210> 83
<211> 540
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 83
Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser
1 5 10 15

Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser Glu Ser Pro Ser
20 25 30

Gly Thr Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro
35 40 45

Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Thr
50 55 60

Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser Glu Ser Pro Ser
65 70 75 80

Gly Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro
85 90 95

Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser
100 105 110

Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser
115 120 125

Ser Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro

130 135 140

Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Pro
145 150 155 160

Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser
165 170 175

Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
180 185 190

Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Thr
195 200 205

Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser Glu Ser Pro Ser
210 215 220

Gly Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
225 230 235 240

Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser
245 250 255

Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Thr Pro Glu Ser Gly
260 265 270

Ser Ala Ser Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
275 280 285

Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Thr
290 295 300

Pro Glu Ser Gly Ser Ala Ser Pro Gly Thr Ser Thr Pro Glu Ser Gly
305 310 315 320

Ser Ala Ser Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
325 330 335

Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser
340 345 350

Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu
355 360 365

Ser Pro Gly Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
370 375 380

Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser
385 390 395 400

Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser

405 410 415

Gly Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
420 425 430

Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser
435 440 445

Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly
450 455 460

Ser Ala Ser Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro
465 470 475 480

Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro
485 490 495

Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu
500 505 510

Ser Pro Gly Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
515 520 525

Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
530 535 540

<210> 84
<211> 576
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 84
Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Gly Gly
1 5 10 15

Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Ser Glu Ser Gly Ser Ser
20 25 30

Glu Gly Gly Pro Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro
35 40 45

Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Ser Glu
50 55 60

Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Ser Glu Ser Gly Ser Ser
65 70 75 80

Glu Gly Gly Pro Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser
85 90 95

```
Gly Ser Glu Gly Ser Ser Gly Pro Gly Glu Ser Ser Gly Ser Ser Glu
            100                 105                 110

Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Ser Glu Ser Gly Ser Ser
        115                 120                 125

Glu Gly Gly Pro Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro
    130                 135                 140

Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Glu Ser Pro
145                 150                 155                 160

Gly Gly Ser Ser Gly Ser Glu Ser Gly Glu Ser Pro Gly Gly Ser Ser
                165                 170                 175

Gly Ser Glu Ser Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser
            180                 185                 190

Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Gly Gly
        195                 200                 205

Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Gly Gly Glu Pro Ser Glu
    210                 215                 220

Ser Gly Ser Ser Gly Ser Glu Gly Ser Ser Gly Pro Gly Glu Ser Ser
225                 230                 235                 240

Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser Gly Ser Gly Gly
                245                 250                 255

Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Gly Gly Glu Pro Ser Glu
            260                 265                 270

Ser Gly Ser Ser Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser
        275                 280                 285

Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro Gly Glu Ser Pro
    290                 295                 300

Gly Gly Ser Ser Gly Ser Glu Ser Gly Glu Ser Pro Gly Gly Ser Ser
305                 310                 315                 320

Gly Ser Glu Ser Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser
                325                 330                 335

Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser Gly Glu Ser Pro
            340                 345                 350

Gly Gly Ser Ser Gly Ser Glu Ser Gly Ser Ser Glu Ser Gly Ser Ser
        355                 360                 365
```

Glu Gly Gly Pro Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser
370 375 380

Gly Ser Glu Gly Ser Ser Gly Pro Gly Glu Ser Ser Gly Ser Ser Glu
385 390 395 400

Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Gly Gly Glu Pro Ser Glu
405 410 415

Ser Gly Ser Ser Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro
420 425 430

Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Glu Ser Pro
435 440 445

Gly Gly Ser Ser Gly Ser Glu Ser Gly Glu Ser Pro Gly Gly Ser Ser
450 455 460

Gly Ser Glu Ser Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro
465 470 475 480

Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Ser Glu
485 490 495

Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Gly Gly Glu Pro Ser Glu
500 505 510

Ser Gly Ser Ser Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser
515 520 525

Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser Gly Ser Glu Gly
530 535 540

Ser Ser Gly Pro Gly Glu Ser Ser Gly Ser Ser Glu Ser Gly Ser Ser
545 550 555 560

Glu Gly Gly Pro Gly Ser Glu Gly Ser Ser Gly Pro Gly Glu Ser Ser
565 570 575

<210> 85
<211> 576
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 85
Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu

```
            20                  25                  30

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
        35                  40                  45

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
    50                  55                  60

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
65                  70                  75                  80

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
                85                  90                  95

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
            100                 105                 110

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
        115                 120                 125

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    130                 135                 140

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
145                 150                 155                 160

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
                165                 170                 175

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            180                 185                 190

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
        195                 200                 205

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
    210                 215                 220

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
225                 230                 235                 240

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
                245                 250                 255

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
            260                 265                 270

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        275                 280                 285

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
```

290 295 300

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
305 310 315 320

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
325 330 335

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
340 345 350

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
355 360 365

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
370 375 380

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
385 390 395 400

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
405 410 415

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
420 425 430

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
435 440 445

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
450 455 460

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
465 470 475 480

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
485 490 495

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
500 505 510

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
515 520 525

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala
530 535 540

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
545 550 555 560

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro

565 570 575

<210> 86
<211> 576
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 86
Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser
1 5 10 15

Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser Glu Ser Pro Ser
20 25 30

Gly Thr Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro
35 40 45

Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Thr
50 55 60

Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser Glu Ser Pro Ser
65 70 75 80

Gly Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro
85 90 95

Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser
100 105 110

Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser
115 120 125

Ser Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
130 135 140

Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Pro
145 150 155 160

Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser
165 170 175

Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
180 185 190

Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Thr
195 200 205

Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser Glu Ser Pro Ser
210 215 220

```
Gly Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
225                 230                 235                 240

Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser
                245                 250                 255

Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Thr Pro Glu Ser Gly
            260                 265                 270

Ser Ala Ser Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
        275                 280                 285

Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Thr
    290                 295                 300

Pro Glu Ser Gly Ser Ala Ser Pro Gly Thr Ser Thr Pro Glu Ser Gly
305                 310                 315                 320

Ser Ala Ser Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
                325                 330                 335

Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser
            340                 345                 350

Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu
        355                 360                 365

Ser Pro Gly Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
    370                 375                 380

Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser
385                 390                 395                 400

Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser
                405                 410                 415

Gly Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
            420                 425                 430

Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser
        435                 440                 445

Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly
    450                 455                 460

Ser Ala Ser Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro
465                 470                 475                 480

Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro
                485                 490                 495
```

Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu
500 505 510

Ser Pro Gly Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
515 520 525

Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser
530 535 540

Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Thr Pro Glu Ser Gly
545 550 555 560

Ser Ala Ser Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
565 570 575

<210> 87
<211> 576
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 87
Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser
1 5 10 15

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
20 25 30

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
35 40 45

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
50 55 60

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
65 70 75 80

Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
85 90 95

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
100 105 110

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser
115 120 125

Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
130 135 140

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser

145 150 155 160

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr
165 170 175

Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
180 185 190

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
195 200 205

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly
210 215 220

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
225 230 235 240

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
245 250 255

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
260 265 270

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
275 280 285

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
290 295 300

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser
305 310 315 320

Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
325 330 335

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser
340 345 350

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
355 360 365

Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
370 375 380

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
385 390 395 400

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
405 410 415

Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser

420 425 430

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro
435 440 445

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
450 455 460

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
465 470 475 480

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser
485 490 495

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser
500 505 510

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
515 520 525

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
530 535 540

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
545 550 555 560

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
565 570 575

<210> 88
<211> 625
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 88
Met Ala Glu Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Pro
1 5 10 15

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
20 25 30

Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
35 40 45

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
50 55 60

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser
65 70 75 80

```
Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
                85                  90                  95

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
            100                 105                 110

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
        115                 120                 125

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
    130                 135                 140

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro
145                 150                 155                 160

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr
                165                 170                 175

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            180                 185                 190

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro
        195                 200                 205

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
    210                 215                 220

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
225                 230                 235                 240

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
                245                 250                 255

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
            260                 265                 270

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
        275                 280                 285

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
    290                 295                 300

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
305                 310                 315                 320

Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
                325                 330                 335

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            340                 345                 350
```

```
Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        355                 360                 365

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    370                 375                 380

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
385                 390                 395                 400

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                405                 410                 415

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            420                 425                 430

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
        435                 440                 445

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro
    450                 455                 460

Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
465                 470                 475                 480

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
                485                 490                 495

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
            500                 505                 510

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
        515                 520                 525

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
    530                 535                 540

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
545                 550                 555                 560

Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
                565                 570                 575

Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro
            580                 585                 590

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr
        595                 600                 605

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
    610                 615                 620
```

Pro
625

<210> 89
<211> 836
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 89
Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Ser Glu
1 5 10 15

Ser Gly Ser Ser Glu Gly Gly Pro Gly Glu Ser Pro Gly Gly Ser Ser
20 25 30

Gly Ser Glu Ser Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser
35 40 45

Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser Gly Glu Ser Pro
50 55 60

Gly Gly Ser Ser Gly Ser Glu Ser Gly Ser Ser Glu Ser Gly Ser Ser
65 70 75 80

Glu Gly Gly Pro Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro
85 90 95

Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro Gly Glu Ser Pro
100 105 110

Gly Gly Ser Ser Gly Ser Glu Ser Gly Glu Ser Pro Gly Gly Ser Ser
115 120 125

Gly Ser Glu Ser Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser
130 135 140

Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Ser Glu
145 150 155 160

Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Ser Glu Ser Gly Ser Ser
165 170 175

Glu Gly Gly Pro Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro
180 185 190

Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Ser Glu
195 200 205

Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Gly Gly Glu Pro Ser Glu

210 215 220

Ser Gly Ser Ser Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser
225 230 235 240

Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser Gly Ser Gly Gly
245 250 255

Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Glu Gly Ser Ser Gly Pro
260 265 270

Gly Glu Ser Ser Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro
275 280 285

Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Glu Gly
290 295 300

Ser Ser Gly Pro Gly Glu Ser Ser Gly Ser Ser Glu Ser Gly Ser Ser
305 310 315 320

Glu Gly Gly Pro Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser
325 330 335

Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser Gly Ser Gly Gly
340 345 350

Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Gly Gly Glu Pro Ser Glu
355 360 365

Ser Gly Ser Ser Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro
370 375 380

Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Gly Gly
385 390 395 400

Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Glu Gly Ser Ser Gly Pro
405 410 415

Gly Glu Ser Ser Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser
420 425 430

Gly Ser Glu Gly Ser Ser Gly Pro Gly Glu Ser Ser Gly Ser Glu Gly
435 440 445

Ser Ser Gly Pro Gly Glu Ser Ser Gly Ser Gly Gly Glu Pro Ser Glu
450 455 460

Ser Gly Ser Ser Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro
465 470 475 480

Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro Gly Glu Ser Pro

485 490 495

Gly Gly Ser Ser Gly Ser Glu Ser Gly Ser Gly Gly Glu Pro Ser Glu
500 505 510

Ser Gly Ser Ser Gly Ser Glu Gly Ser Ser Gly Pro Gly Glu Ser Ser
515 520 525

Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser Gly Ser Glu Gly
530 535 540

Ser Ser Gly Pro Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro
545 550 555 560

Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Glu Gly
565 570 575

Ser Ser Gly Pro Gly Glu Ser Ser Gly Ser Glu Gly Ser Ser Gly Pro
580 585 590

Gly Glu Ser Ser Gly Ser Glu Gly Ser Ser Gly Pro Gly Glu Ser Ser
595 600 605

Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Gly Gly
610 615 620

Glu Pro Ser Glu Ser Gly Ser Ser Gly Glu Ser Pro Gly Gly Ser Ser
625 630 635 640

Gly Ser Glu Ser Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser
645 650 655

Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Glu Gly
660 665 670

Ser Ser Gly Pro Gly Glu Ser Ser Gly Glu Ser Pro Gly Gly Ser Ser
675 680 685

Gly Ser Glu Ser Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro
690 695 700

Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Ser Glu
705 710 715 720

Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Gly Gly Glu Pro Ser Glu
725 730 735

Ser Gly Ser Ser Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro
740 745 750

Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser Gly Ser Gly Gly

755 760 765

Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Ser Glu Ser Gly Ser Ser
770 775 780

Glu Gly Gly Pro Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser
785 790 795 800

Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Glu Ser Pro
805 810 815

Gly Gly Ser Ser Gly Ser Glu Ser Gly Ser Gly Gly Glu Pro Ser Glu
820 825 830

Ser Gly Ser Ser
835

<210> 90
<211> 864
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 90
Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
35 40 45

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
50 55 60

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
65 70 75 80

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
85 90 95

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
100 105 110

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
115 120 125

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
130 135 140

```
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
145                 150                 155                 160

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
                165                 170                 175

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            180                 185                 190

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
        195                 200                 205

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
    210                 215                 220

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
225                 230                 235                 240

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
                245                 250                 255

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
            260                 265                 270

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        275                 280                 285

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
    290                 295                 300

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
305                 310                 315                 320

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
                325                 330                 335

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
            340                 345                 350

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
        355                 360                 365

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    370                 375                 380

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
385                 390                 395                 400

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
                405                 410                 415
```

```
Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            420                 425                 430

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
        435                 440                 445

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
    450                 455                 460

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
465                 470                 475                 480

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
                485                 490                 495

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
            500                 505                 510

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
        515                 520                 525

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala
    530                 535                 540

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
545                 550                 555                 560

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
                565                 570                 575

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
            580                 585                 590

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
        595                 600                 605

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
    610                 615                 620

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
625                 630                 635                 640

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
                645                 650                 655

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
            660                 665                 670

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
        675                 680                 685
```

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
690 695 700

Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
705 710 715 720

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
725 730 735

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
740 745 750

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
755 760 765

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
770 775 780

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
785 790 795 800

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
805 810 815

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
820 825 830

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
835 840 845

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
850 855 860

<210> 91
<211> 875
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<220>
<221> MOD_RES
<222> (430)..(432)
<223> Any amino acid

<220>
<221> MOD_RES
<222> (443)..(446)
<223> Any amino acid

<400> 91
Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Pro
1 5 10 15

```
Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser
            20                  25                  30

Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
        35                  40                  45

Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Thr Ser Thr
    50                  55                  60

Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser Glu Ser Pro Ser
65                  70                  75                  80

Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
                85                  90                  95

Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser
            100                 105                 110

Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser
        115                 120                 125

Ser Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro
    130                 135                 140

Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro
145                 150                 155                 160

Ser Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser
                165                 170                 175

Ser Thr Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro
            180                 185                 190

Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Thr Ser Thr
        195                 200                 205

Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser Glu Ser Pro Ser
    210                 215                 220

Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
225                 230                 235                 240

Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser
                245                 250                 255

Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Thr Pro Glu Ser Gly
            260                 265                 270

Ser Ala Ser Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
        275                 280                 285
```

```
Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser
    290                 295                 300

Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro Ser Gly Glu Ser
305                 310                 315                 320

Ser Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
                325                 330                 335

Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser
            340                 345                 350

Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser Ser Thr Ala Glu
        355                 360                 365

Ser Pro Gly Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro
    370                 375                 380

Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser
385                 390                 395                 400

Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser
                405                 410                 415

Gly Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Pro Xaa Xaa Xaa
            420                 425                 430

Gly Ala Ser Ala Ser Gly Ala Pro Ser Thr Xaa Xaa Xaa Xaa Ser Glu
        435                 440                 445

Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly
    450                 455                 460

Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly
465                 470                 475                 480

Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu
                485                 490                 495

Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly
            500                 505                 510

Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly
        515                 520                 525

Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Pro Ser
    530                 535                 540

Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser
545                 550                 555                 560
```

```
Pro Gly Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly
                565                 570                 575

Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser Glu
            580                 585                 590

Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly
        595                 600                 605

Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly
    610                 615                 620

Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Thr Pro
625                 630                 635                 640

Glu Ser Gly Ser Ala Ser Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser
                645                 650                 655

Ala Ser Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly
            660                 665                 670

Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser Ser
        675                 680                 685

Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly
    690                 695                 700

Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly
705                 710                 715                 720

Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser Ser
                725                 730                 735

Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser
            740                 745                 750

Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly
        755                 760                 765

Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Pro Ser
    770                 775                 780

Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser
785                 790                 795                 800

Thr Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly
                805                 810                 815

Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro Ser
            820                 825                 830
```

Gly Glu Ser Ser Thr Ala Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
835 840 845

Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
850 855 860

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
865 870 875

<210> 92
<211> 864
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 92
Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
1 5 10 15

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
20 25 30

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
35 40 45

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
50 55 60

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
65 70 75 80

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
85 90 95

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
100 105 110

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
115 120 125

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
130 135 140

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
145 150 155 160

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
165 170 175

```
Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
            180                 185                 190

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
        195                 200                 205

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
    210                 215                 220

Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
225                 230                 235                 240

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro
                245                 250                 255

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
            260                 265                 270

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
        275                 280                 285

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
    290                 295                 300

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
305                 310                 315                 320

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                325                 330                 335

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
            340                 345                 350

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
        355                 360                 365

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    370                 375                 380

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
385                 390                 395                 400

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
                405                 410                 415

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
            420                 425                 430

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
        435                 440                 445
```

```
Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
    450                 455                 460

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
465                 470                 475                 480

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
                485                 490                 495

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
            500                 505                 510

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
        515                 520                 525

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
    530                 535                 540

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
545                 550                 555                 560

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                565                 570                 575

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
            580                 585                 590

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
        595                 600                 605

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
    610                 615                 620

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
625                 630                 635                 640

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
                645                 650                 655

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            660                 665                 670

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
        675                 680                 685

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
    690                 695                 700

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
705                 710                 715                 720
```

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
725 730 735

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
740 745 750

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
755 760 765

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
770 775 780

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
785 790 795 800

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
805 810 815

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
820 825 830

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
835 840 845

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
850 855 860

<210> 93
<211> 875
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 93
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
1 5 10 15

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
20 25 30

Ser Thr Glu Glu Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro
35 40 45

Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser
50 55 60

Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser
65 70 75 80

Gly Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
85 90 95

```
Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Glu Pro
            100                 105                 110

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
        115                 120                 125

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
    130                 135                 140

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
145                 150                 155                 160

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
                165                 170                 175

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            180                 185                 190

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
        195                 200                 205

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
    210                 215                 220

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
225                 230                 235                 240

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
                245                 250                 255

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
            260                 265                 270

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        275                 280                 285

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
    290                 295                 300

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
305                 310                 315                 320

Ser Thr Glu Glu Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
                325                 330                 335

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
            340                 345                 350

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Ser Thr Glu Pro Ser Glu
        355                 360                 365
```

```
Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    370                 375                 380

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
385                 390                 395                 400

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
                405                 410                 415

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            420                 425                 430

Gly Ala Ser Ala Ser Gly Ala Pro Ser Thr Gly Gly Thr Ser Glu Ser
        435                 440                 445

Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
    450                 455                 460

Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
465                 470                 475                 480

Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser Glu
                485                 490                 495

Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser
            500                 505                 510

Thr Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
        515                 520                 525

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser
    530                 535                 540

Ala Ser Thr Gly Thr Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
545                 550                 555                 560

Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
                565                 570                 575

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Thr Ser Ser
            580                 585                 590

Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser
        595                 600                 605

Pro Gly Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly
    610                 615                 620

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala
625                 630                 635                 640
```

```
Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                645                 650                 655

Ser Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly
            660                 665                 670

Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser Glu
        675                 680                 685

Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    690                 695                 700

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
705                 710                 715                 720

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Ser Thr Pro
                725                 730                 735

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
            740                 745                 750

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
        755                 760                 765

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
    770                 775                 780

Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
785                 790                 795                 800

Thr Glu Glu Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
                805                 810                 815

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly
            820                 825                 830

Thr Ser Ser Thr Gly Ser Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
        835                 840                 845

Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
    850                 855                 860

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
865                 870                 875


<210> 94
<211> 913
<212> PRT
<213> Artificial Sequence

<220>
```

<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 94
Met Ala Glu Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Pro
1 5 10 15

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
20 25 30

Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
35 40 45

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
50 55 60

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser
65 70 75 80

Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
85 90 95

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
100 105 110

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
115 120 125

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
130 135 140

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro
145 150 155 160

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr
165 170 175

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
180 185 190

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro
195 200 205

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
210 215 220

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
225 230 235 240

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
245 250 255

```
Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
            260                 265                 270

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
        275                 280                 285

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
    290                 295                 300

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
305                 310                 315                 320

Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
                325                 330                 335

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            340                 345                 350

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        355                 360                 365

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    370                 375                 380

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
385                 390                 395                 400

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                405                 410                 415

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            420                 425                 430

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
        435                 440                 445

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro
    450                 455                 460

Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
465                 470                 475                 480

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
                485                 490                 495

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
            500                 505                 510

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
        515                 520                 525
```

```
Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
    530                 535                 540

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
545                 550                 555                 560

Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
                565                 570                 575

Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro
            580                 585                 590

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr
        595                 600                 605

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
    610                 615                 620

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
625                 630                 635                 640

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
                645                 650                 655

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
            660                 665                 670

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
        675                 680                 685

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro
    690                 695                 700

Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
705                 710                 715                 720

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser
                725                 730                 735

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro
            740                 745                 750

Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
        755                 760                 765

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
    770                 775                 780

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
785                 790                 795                 800
```

Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
805 810 815

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu
820 825 830

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro
835 840 845

Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
850 855 860

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu
865 870 875 880

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
885 890 895

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
900 905 910

Pro

<210> 95
<211> 924
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 95
Met Ala Glu Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ala Ser
1 5 10 15

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly
20 25 30

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
35 40 45

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu
50 55 60

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro
65 70 75 80

Thr Ser Thr Glu Glu Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly
85 90 95

Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr
100 105 110

```
Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro
        115                 120                 125

Ser Gly Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser
    130                 135                 140

Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Glu
145                 150                 155                 160

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
                165                 170                 175

Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
            180                 185                 190

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
        195                 200                 205

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser
    210                 215                 220

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
225                 230                 235                 240

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
                245                 250                 255

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
            260                 265                 270

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
        275                 280                 285

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
    290                 295                 300

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
305                 310                 315                 320

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
                325                 330                 335

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu
            340                 345                 350

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro
        355                 360                 365

Thr Ser Thr Glu Glu Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
    370                 375                 380
```

```
Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser
385                 390                 395                 400

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Ser Thr Glu Pro Ser
                405                 410                 415

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            420                 425                 430

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro
        435                 440                 445

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro
    450                 455                 460

Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
465                 470                 475                 480

Pro Gly Ala Ser Ala Ser Gly Ala Pro Ser Thr Gly Gly Thr Ser Glu
                485                 490                 495

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
            500                 505                 510

Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
        515                 520                 525

Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser
    530                 535                 540

Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser
545                 550                 555                 560

Ser Thr Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
                565                 570                 575

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
            580                 585                 590

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
        595                 600                 605

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
    610                 615                 620

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Thr Ser
625                 630                 635                 640

Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser Ser Thr Ala Glu
                645                 650                 655
```

```
Ser Pro Gly Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro
            660                 665                 670

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
        675                 680                 685

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu
    690                 695                 700

Gly Ser Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro
705                 710                 715                 720

Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser
                725                 730                 735

Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
            740                 745                 750

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
        755                 760                 765

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Ser Thr
    770                 775                 780

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
785                 790                 795                 800

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                805                 810                 815

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
            820                 825                 830

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
        835                 840                 845

Ser Thr Glu Glu Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
    850                 855                 860

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
865                 870                 875                 880

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Ser Glu Ser Ala Thr Pro
                885                 890                 895

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            900                 905                 910

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
        915                 920
```

<210> 96
<211> 1318
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 96
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
1 5 10 15

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
20 25 30

Ser Thr Glu Glu Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro
35 40 45

Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser
50 55 60

Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser
65 70 75 80

Gly Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
85 90 95

Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Glu Pro
100 105 110

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
115 120 125

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
130 135 140

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
145 150 155 160

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
165 170 175

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
180 185 190

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
195 200 205

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
210 215 220

```
Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
225                 230                 235                 240

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
                245                 250                 255

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
            260                 265                 270

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        275                 280                 285

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
    290                 295                 300

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
305                 310                 315                 320

Ser Thr Glu Glu Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
                325                 330                 335

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
            340                 345                 350

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Ser Thr Glu Pro Ser Glu
        355                 360                 365

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    370                 375                 380

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
385                 390                 395                 400

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
                405                 410                 415

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            420                 425                 430

Gly Pro Glu Pro Thr Gly Pro Ala Pro Ser Gly Gly Ser Glu Pro Ala
        435                 440                 445

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
    450                 455                 460

Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
465                 470                 475                 480

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly
                485                 490                 495
```

```
Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser
            500                 505                 510

Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
        515                 520                 525

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
    530                 535                 540

Ser Pro Thr Ser Thr Glu Glu Gly Ser Thr Ser Ser Thr Ala Glu Ser
545                 550                 555                 560

Pro Gly Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly
                565                 570                 575

Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser Glu
            580                 585                 590

Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly
        595                 600                 605

Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly
    610                 615                 620

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
625                 630                 635                 640

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
                645                 650                 655

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
            660                 665                 670

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser
        675                 680                 685

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    690                 695                 700

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
705                 710                 715                 720

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Pro Ser
                725                 730                 735

Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser
            740                 745                 750

Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly
        755                 760                 765
```

```
Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
    770                 775                 780

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
785                 790                 795                 800

Ser Ala Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
                805                 810                 815

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
            820                 825                 830

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
        835                 840                 845

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
    850                 855                 860

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Ala Ser
865                 870                 875                 880

Gly Ala Pro Ser Thr Gly Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr
                885                 890                 895

Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr
            900                 905                 910

Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Glu Ser Ala
        915                 920                 925

Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
    930                 935                 940

Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
945                 950                 955                 960

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser
                965                 970                 975

Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
            980                 985                 990

Ser Pro Gly Thr Ser Thr Pro Glu  Ser Gly Ser Ala Ser  Pro Gly Thr
        995                 1000                1005

Ser Pro  Ser Gly Glu Ser Ser  Thr Ala Pro Gly Thr  Ser Pro Ser
    1010                1015                1020

Gly Glu  Ser Ser Thr Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1025                1030                1035
```

```
Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1040                1045                1050

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    1055                1060                1065

Thr Ser  Glu Ser Pro Ser Gly  Thr Ala Pro Gly Ser  Thr Ser Glu
    1070                1075                1080

Ser Pro  Ser Gly Thr Ala Pro  Gly Thr Ser Thr Pro  Glu Ser Gly
    1085                1090                1095

Ser Ala  Ser Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    1100                1105                1110

Glu Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1115                1120                1125

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1130                1135                1140

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1145                1150                1155

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1160                1165                1170

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ala
    1175                1180                1185

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ser  Ser Thr Pro
    1190                1195                1200

Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Thr Ser Glu  Ser Pro Ser
    1205                1210                1215

Gly Thr  Ala Pro Gly Thr Ser  Pro Ser Gly Glu Ser  Ser Thr Ala
    1220                1225                1230

Pro Gly  Ser Thr Ser Ser Thr  Ala Glu Ser Pro Gly  Pro Gly Ser
    1235                1240                1245

Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1250                1255                1260

Thr Ser  Ser Thr Gly Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala
    1265                1270                1275

Ser Ser  Ser Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    1280                1285                1290
```

Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr
1295 1300 1305

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
1310 1315

<210> 97
<211> 864
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 97
Gly Thr Ser Thr Glu Pro Ser Glu Pro Gly Ser Ala Gly Thr Ser Thr
1 5 10 15

Glu Pro Ser Glu Pro Gly Ser Ala Gly Ser Glu Pro Ala Thr Ser Gly
20 25 30

Thr Glu Pro Ser Gly Ser Gly Ala Ser Glu Pro Thr Ser Thr Glu Pro
35 40 45

Gly Ser Glu Pro Ala Thr Ser Gly Thr Glu Pro Ser Gly Ser Glu Pro
50 55 60

Ala Thr Ser Gly Thr Glu Pro Ser Gly Ser Glu Pro Ala Thr Ser Gly
65 70 75 80

Thr Glu Pro Ser Gly Ser Gly Ala Ser Glu Pro Thr Ser Thr Glu Pro
85 90 95

Gly Thr Ser Thr Glu Pro Ser Glu Pro Gly Ser Ala Gly Ser Glu Pro
100 105 110

Ala Thr Ser Gly Thr Glu Pro Ser Gly Thr Ser Thr Glu Pro Ser Glu
115 120 125

Pro Gly Ser Ala Gly Ser Glu Pro Ala Thr Ser Gly Thr Glu Pro Ser
130 135 140

Gly Ser Glu Pro Ala Thr Ser Gly Thr Glu Pro Ser Gly Thr Ser Thr
145 150 155 160

Glu Pro Ser Glu Pro Gly Ser Ala Gly Thr Ser Thr Glu Pro Ser Glu
165 170 175

Pro Gly Ser Ala Gly Ser Glu Pro Ala Thr Ser Gly Thr Glu Pro Ser
180 185 190

Gly Ser Glu Pro Ala Thr Ser Gly Thr Glu Pro Ser Gly Thr Ser Glu
195 200 205

```
Pro Ser Thr Ser Glu Pro Gly Ala Gly Ser Gly Ala Ser Glu Pro Thr
    210                 215                 220

Ser Thr Glu Pro Gly Thr Ser Glu Pro Ser Thr Ser Glu Pro Gly Ala
225                 230                 235                 240

Gly Ser Glu Pro Ala Thr Ser Gly Thr Glu Pro Ser Gly Ser Glu Pro
                245                 250                 255

Ala Thr Ser Gly Thr Glu Pro Ser Gly Thr Ser Thr Glu Pro Ser Glu
            260                 265                 270

Pro Gly Ser Ala Gly Thr Ser Thr Glu Pro Ser Glu Pro Gly Ser Ala
        275                 280                 285

Gly Ser Gly Ala Ser Glu Pro Thr Ser Thr Glu Pro Gly Ser Glu Pro
    290                 295                 300

Ala Thr Ser Gly Thr Glu Pro Ser Gly Ser Glu Pro Ala Thr Ser Gly
305                 310                 315                 320

Thr Glu Pro Ser Gly Ser Glu Pro Ala Thr Ser Gly Thr Glu Pro Ser
                325                 330                 335

Gly Ser Glu Pro Ala Thr Ser Gly Thr Glu Pro Ser Gly Thr Ser Thr
            340                 345                 350

Glu Pro Ser Glu Pro Gly Ser Ala Gly Ser Glu Pro Ala Thr Ser Gly
        355                 360                 365

Thr Glu Pro Ser Gly Ser Gly Ala Ser Glu Pro Thr Ser Thr Glu Pro
    370                 375                 380

Gly Thr Ser Thr Glu Pro Ser Glu Pro Gly Ser Ala Gly Ser Glu Pro
385                 390                 395                 400

Ala Thr Ser Gly Thr Glu Pro Ser Gly Ser Gly Ala Ser Glu Pro Thr
                405                 410                 415

Ser Thr Glu Pro Gly Thr Ser Thr Glu Pro Ser Glu Pro Gly Ser Ala
            420                 425                 430

Gly Ser Gly Ala Ser Glu Pro Thr Ser Thr Glu Pro Gly Ser Glu Pro
        435                 440                 445

Ala Thr Ser Gly Thr Glu Pro Ser Gly Ser Gly Ala Ser Glu Pro Thr
    450                 455                 460

Ser Thr Glu Pro Gly Ser Glu Pro Ala Thr Ser Gly Thr Glu Pro Ser
465                 470                 475                 480
```

```
Gly Ser Gly Ala Ser Glu Pro Thr Ser Thr Glu Pro Gly Thr Ser Thr
                485                 490                 495

Glu Pro Ser Glu Pro Gly Ser Ala Gly Ser Glu Pro Ala Thr Ser Gly
            500                 505                 510

Thr Glu Pro Ser Gly Ser Gly Ala Ser Glu Pro Thr Ser Thr Glu Pro
        515                 520                 525

Gly Thr Ser Thr Glu Pro Ser Glu Pro Gly Ser Ala Gly Ser Glu Pro
    530                 535                 540

Ala Thr Ser Gly Thr Glu Pro Ser Gly Thr Ser Thr Glu Pro Ser Glu
545                 550                 555                 560

Pro Gly Ser Ala Gly Ser Glu Pro Ala Thr Ser Gly Thr Glu Pro Ser
                565                 570                 575

Gly Thr Ser Thr Glu Pro Ser Glu Pro Gly Ser Ala Gly Thr Ser Thr
            580                 585                 590

Glu Pro Ser Glu Pro Gly Ser Ala Gly Thr Ser Thr Glu Pro Ser Glu
        595                 600                 605

Pro Gly Ser Ala Gly Thr Ser Thr Glu Pro Ser Glu Pro Gly Ser Ala
    610                 615                 620

Gly Thr Ser Thr Glu Pro Ser Glu Pro Gly Ser Ala Gly Thr Ser Thr
625                 630                 635                 640

Glu Pro Ser Glu Pro Gly Ser Ala Gly Thr Ser Glu Pro Ser Thr Ser
                645                 650                 655

Glu Pro Gly Ala Gly Ser Gly Ala Ser Glu Pro Thr Ser Thr Glu Pro
            660                 665                 670

Gly Thr Ser Thr Glu Pro Ser Glu Pro Gly Ser Ala Gly Thr Ser Thr
        675                 680                 685

Glu Pro Ser Glu Pro Gly Ser Ala Gly Thr Ser Thr Glu Pro Ser Glu
    690                 695                 700

Pro Gly Ser Ala Gly Ser Glu Pro Ala Thr Ser Gly Thr Glu Pro Ser
705                 710                 715                 720

Gly Ser Gly Ala Ser Glu Pro Thr Ser Thr Glu Pro Gly Ser Glu Pro
                725                 730                 735

Ala Thr Ser Gly Thr Glu Pro Ser Gly Ser Glu Pro Ala Thr Ser Gly
            740                 745                 750
```

Thr Glu Pro Ser Gly Ser Glu Pro Ala Thr Ser Gly Thr Glu Pro Ser
755 760 765

Gly Ser Glu Pro Ala Thr Ser Gly Thr Glu Pro Ser Gly Thr Ser Glu
770 775 780

Pro Ser Thr Ser Glu Pro Gly Ala Gly Ser Glu Pro Ala Thr Ser Gly
785 790 795 800

Thr Glu Pro Ser Gly Ser Gly Ala Ser Glu Pro Thr Ser Thr Glu Pro
805 810 815

Gly Thr Ser Thr Glu Pro Ser Glu Pro Gly Ser Ala Gly Ser Glu Pro
820 825 830

Ala Thr Ser Gly Thr Glu Pro Ser Gly Ser Gly Ala Ser Glu Pro Thr
835 840 845

Ser Thr Glu Pro Gly Thr Ser Thr Glu Pro Ser Glu Pro Gly Ser Ala
850 855 860

<210> 98
<211> 864
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 98
Gly Ser Glu Thr Ala Thr Ser Gly Ser Glu Thr Ala Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Ser Glu Ser Gly Ala Gly Ser Thr Ala Gly Ser Glu Thr
20 25 30

Ser Thr Glu Ala Gly Thr Ser Glu Ser Ala Thr Ser Glu Ser Gly Ala
35 40 45

Gly Ser Glu Thr Ala Thr Ser Gly Ser Glu Thr Ala Gly Ser Glu Thr
50 55 60

Ala Thr Ser Gly Ser Glu Thr Ala Gly Thr Ser Thr Glu Ala Ser Glu
65 70 75 80

Gly Ser Ala Ser Gly Thr Ser Thr Glu Ala Ser Glu Gly Ser Ala Ser
85 90 95

Gly Thr Ser Glu Ser Ala Thr Ser Glu Ser Gly Ala Gly Ser Glu Thr
100 105 110

```
Ala Thr Ser Gly Ser Glu Thr Ala Gly Thr Ser Thr Glu Ala Ser Glu
        115                 120                 125

Gly Ser Ala Ser Gly Ser Thr Ala Gly Ser Glu Thr Ser Thr Glu Ala
    130                 135                 140

Gly Thr Ser Glu Ser Ala Thr Ser Glu Ser Gly Ala Gly Thr Ser Glu
145                 150                 155                 160

Ser Ala Thr Ser Glu Ser Gly Ala Gly Ser Glu Thr Ala Thr Ser Gly
                165                 170                 175

Ser Glu Thr Ala Gly Thr Ser Glu Ser Ala Thr Ser Glu Ser Gly Ala
            180                 185                 190

Gly Thr Ser Thr Glu Ala Ser Glu Gly Ser Ala Ser Gly Ser Glu Thr
        195                 200                 205

Ala Thr Ser Gly Ser Glu Thr Ala Gly Ser Glu Thr Ala Thr Ser Gly
    210                 215                 220

Ser Glu Thr Ala Gly Thr Ser Thr Glu Ala Ser Glu Gly Ser Ala Ser
225                 230                 235                 240

Gly Ser Thr Ala Gly Ser Glu Thr Ser Thr Glu Ala Gly Thr Ser Glu
                245                 250                 255

Ser Ala Thr Ser Glu Ser Gly Ala Gly Thr Ser Thr Glu Ala Ser Glu
            260                 265                 270

Gly Ser Ala Ser Gly Ser Glu Thr Ala Thr Ser Gly Ser Glu Thr Ala
        275                 280                 285

Gly Ser Thr Ala Gly Ser Glu Thr Ser Thr Glu Ala Gly Ser Thr Ala
    290                 295                 300

Gly Ser Glu Thr Ser Thr Glu Ala Gly Ser Glu Thr Ala Thr Ser Gly
305                 310                 315                 320

Ser Glu Thr Ala Gly Thr Ser Glu Ser Ala Thr Ser Glu Ser Gly Ala
                325                 330                 335

Gly Thr Ser Glu Ser Ala Thr Ser Glu Ser Gly Ala Gly Ser Glu Thr
            340                 345                 350

Ala Thr Ser Gly Ser Glu Thr Ala Gly Thr Ser Glu Ser Ala Thr Ser
        355                 360                 365

Glu Ser Gly Ala Gly Thr Ser Glu Ser Ala Thr Ser Glu Ser Gly Ala
    370                 375                 380
```

```
Gly Ser Glu Thr Ala Thr Ser Gly Ser Glu Thr Ala Gly Ser Glu Thr
385                 390                 395                 400

Ala Thr Ser Gly Ser Glu Thr Ala Gly Thr Ser Thr Glu Ala Ser Glu
                405                 410                 415

Gly Ser Ala Ser Gly Ser Thr Ala Gly Ser Glu Thr Ser Thr Glu Ala
            420                 425                 430

Gly Ser Glu Thr Ala Thr Ser Gly Ser Glu Thr Ala Gly Thr Ser Glu
        435                 440                 445

Ser Ala Thr Ser Glu Ser Gly Ala Gly Ser Thr Ala Gly Ser Glu Thr
    450                 455                 460

Ser Thr Glu Ala Gly Ser Thr Ala Gly Ser Glu Thr Ser Thr Glu Ala
465                 470                 475                 480

Gly Ser Thr Ala Gly Ser Glu Thr Ser Thr Glu Ala Gly Thr Ser Thr
                485                 490                 495

Glu Ala Ser Glu Gly Ser Ala Ser Gly Ser Thr Ala Gly Ser Glu Thr
            500                 505                 510

Ser Thr Glu Ala Gly Ser Thr Ala Gly Ser Glu Thr Ser Thr Glu Ala
        515                 520                 525

Gly Thr Ser Thr Glu Ala Ser Glu Gly Ser Ala Ser Gly Ser Thr Ala
    530                 535                 540

Gly Ser Glu Thr Ser Thr Glu Ala Gly Ser Glu Thr Ala Thr Ser Gly
545                 550                 555                 560

Ser Glu Thr Ala Gly Thr Ser Thr Glu Ala Ser Glu Gly Ser Ala Ser
                565                 570                 575

Gly Thr Ser Glu Ser Ala Thr Ser Glu Ser Gly Ala Gly Ser Glu Thr
            580                 585                 590

Ala Thr Ser Gly Ser Glu Thr Ala Gly Thr Ser Glu Ser Ala Thr Ser
        595                 600                 605

Glu Ser Gly Ala Gly Thr Ser Glu Ser Ala Thr Ser Glu Ser Gly Ala
    610                 615                 620

Gly Ser Glu Thr Ala Thr Ser Gly Ser Glu Thr Ala Gly Thr Ser Glu
625                 630                 635                 640

Ser Ala Thr Ser Glu Ser Gly Ala Gly Ser Glu Thr Ala Thr Ser Gly
                645                 650                 655
```

Ser Glu Thr Ala Gly Thr Ser Thr Glu Ala Ser Glu Gly Ser Ala Ser
660 665 670

Gly Thr Ser Thr Glu Ala Ser Glu Gly Ser Ala Ser Gly Ser Thr Ala
675 680 685

Gly Ser Glu Thr Ser Thr Glu Ala Gly Ser Thr Ala Gly Ser Glu Thr
690 695 700

Ser Thr Glu Ala Gly Ser Glu Thr Ala Thr Ser Gly Ser Glu Thr Ala
705 710 715 720

Gly Thr Ser Glu Ser Ala Thr Ser Glu Ser Gly Ala Gly Thr Ser Glu
725 730 735

Ser Ala Thr Ser Glu Ser Gly Ala Gly Ser Glu Thr Ala Thr Ser Gly
740 745 750

Ser Glu Thr Ala Gly Ser Glu Thr Ala Thr Ser Gly Ser Glu Thr Ala
755 760 765

Gly Ser Glu Thr Ala Thr Ser Gly Ser Glu Thr Ala Gly Thr Ser Thr
770 775 780

Glu Ala Ser Glu Gly Ser Ala Ser Gly Thr Ser Glu Ser Ala Thr Ser
785 790 795 800

Glu Ser Gly Ala Gly Ser Glu Thr Ala Thr Ser Gly Ser Glu Thr Ala
805 810 815

Gly Ser Glu Thr Ala Thr Ser Gly Ser Glu Thr Ala Gly Thr Ser Glu
820 825 830

Ser Ala Thr Ser Glu Ser Gly Ala Gly Thr Ser Glu Ser Ala Thr Ser
835 840 845

Glu Ser Gly Ala Gly Ser Glu Thr Ala Thr Ser Gly Ser Glu Thr Ala
850 855 860

<210> 99
<211> 948
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 99
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
20 25 30

```
Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        35                  40                  45

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
    50                  55                  60

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
65                  70                  75                  80

Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
                85                  90                  95

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu
            100                 105                 110

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
        115                 120                 125

Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    130                 135                 140

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
145                 150                 155                 160

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
                165                 170                 175

Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
            180                 185                 190

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
        195                 200                 205

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
    210                 215                 220

Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
225                 230                 235                 240

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
                245                 250                 255

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu
            260                 265                 270

Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
        275                 280                 285

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
    290                 295                 300
```

```
Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
305                 310                 315                 320

Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
                325                 330                 335

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
            340                 345                 350

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
        355                 360                 365

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
    370                 375                 380

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
385                 390                 395                 400

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
                405                 410                 415

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            420                 425                 430

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
        435                 440                 445

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
    450                 455                 460

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
465                 470                 475                 480

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
                485                 490                 495

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
            500                 505                 510

Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
        515                 520                 525

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu
    530                 535                 540

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
545                 550                 555                 560

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
                565                 570                 575
```

```
Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
            580                 585                 590

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
        595                 600                 605

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
    610                 615                 620

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
625                 630                 635                 640

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
                645                 650                 655

Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
            660                 665                 670

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
        675                 680                 685

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
    690                 695                 700

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
705                 710                 715                 720

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
                725                 730                 735

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
            740                 745                 750

Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
        755                 760                 765

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
    770                 775                 780

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu
785                 790                 795                 800

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
                805                 810                 815

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala
            820                 825                 830

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
        835                 840                 845
```

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
850 855 860

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu
865 870 875 880

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
885 890 895

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
900 905 910

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
915 920 925

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
930 935 940

Glu Ser Gly Pro
945

<210> 100
<211> 1024
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 100
Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
1 5 10 15

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
35 40 45

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
50 55 60

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
65 70 75 80

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
85 90 95

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
100 105 110

```
Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
        115                 120                 125

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
    130                 135                 140

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
145                 150                 155                 160

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
                165                 170                 175

Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            180                 185                 190

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala
        195                 200                 205

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
    210                 215                 220

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
225                 230                 235                 240

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
                245                 250                 255

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
            260                 265                 270

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        275                 280                 285

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
    290                 295                 300

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
305                 310                 315                 320

Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
                325                 330                 335

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
            340                 345                 350

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
        355                 360                 365

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    370                 375                 380
```

```
Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
385                 390                 395                 400

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
                405                 410                 415

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
            420                 425                 430

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
        435                 440                 445

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu
    450                 455                 460

Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
465                 470                 475                 480

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
                485                 490                 495

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
            500                 505                 510

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
        515                 520                 525

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
    530                 535                 540

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
545                 550                 555                 560

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
                565                 570                 575

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
            580                 585                 590

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
        595                 600                 605

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
    610                 615                 620

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
625                 630                 635                 640

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
                645                 650                 655
```

```
Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
            660                 665                 670

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu
        675                 680                 685

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
    690                 695                 700

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
705                 710                 715                 720

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala
                725                 730                 735

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
            740                 745                 750

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
        755                 760                 765

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
    770                 775                 780

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
785                 790                 795                 800

Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
                805                 810                 815

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
            820                 825                 830

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
        835                 840                 845

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
    850                 855                 860

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr
865                 870                 875                 880

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
                885                 890                 895

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
            900                 905                 910

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
        915                 920                 925
```

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
930 935 940

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
945 950 955 960

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
965 970 975

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
980 985 990

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
995 1000 1005

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
1010 1015 1020

Thr

<210> 101
<211> 1024
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 101
Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
1 5 10 15

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
35 40 45

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
50 55 60

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
65 70 75 80

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
85 90 95

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr
100 105 110

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
115 120 125

```
Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
    130                 135                 140

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
145                 150                 155                 160

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
                165                 170                 175

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
            180                 185                 190

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
        195                 200                 205

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
    210                 215                 220

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
225                 230                 235                 240

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
                245                 250                 255

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
            260                 265                 270

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        275                 280                 285

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
    290                 295                 300

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
305                 310                 315                 320

Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
                325                 330                 335

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
            340                 345                 350

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
        355                 360                 365

Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
    370                 375                 380

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
385                 390                 395                 400
```

```
Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
                405                 410                 415

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
            420                 425                 430

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala
        435                 440                 445

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
    450                 455                 460

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
465                 470                 475                 480

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
                485                 490                 495

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
            500                 505                 510

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        515                 520                 525

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
    530                 535                 540

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
545                 550                 555                 560

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
                565                 570                 575

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
            580                 585                 590

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
        595                 600                 605

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
    610                 615                 620

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
625                 630                 635                 640

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
                645                 650                 655

Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            660                 665                 670
```

```
Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
        675                 680                 685

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
    690                 695                 700

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
705                 710                 715                 720

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
                725                 730                 735

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
            740                 745                 750

Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
        755                 760                 765

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
    770                 775                 780

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
785                 790                 795                 800

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
                805                 810                 815

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
            820                 825                 830

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
        835                 840                 845

Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
    850                 855                 860

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
865                 870                 875                 880

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
                885                 890                 895

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
            900                 905                 910

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
        915                 920                 925

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
    930                 935                 940
```

Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
945 950 955 960

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
965 970 975

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
980 985 990

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
995 1000 1005

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro
1010 1015 1020

Ala

<210> 102
<211> 1024
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 102
Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
1 5 10 15

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
20 25 30

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
35 40 45

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr
50 55 60

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
65 70 75 80

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
85 90 95

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
100 105 110

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
115 120 125

```
Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
    130                 135                 140

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
145                 150                 155                 160

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
                165                 170                 175

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
            180                 185                 190

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
        195                 200                 205

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
    210                 215                 220

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
225                 230                 235                 240

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
                245                 250                 255

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
            260                 265                 270

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
        275                 280                 285

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
    290                 295                 300

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
305                 310                 315                 320

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
                325                 330                 335

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
            340                 345                 350

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
        355                 360                 365

Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
    370                 375                 380

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu
385                 390                 395                 400
```

```
Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
                405                 410                 415

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            420                 425                 430

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
        435                 440                 445

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
    450                 455                 460

Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
465                 470                 475                 480

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
                485                 490                 495

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
            500                 505                 510

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
        515                 520                 525

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala
    530                 535                 540

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
545                 550                 555                 560

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
                565                 570                 575

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu
            580                 585                 590

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
        595                 600                 605

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
    610                 615                 620

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
625                 630                 635                 640

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
                645                 650                 655

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            660                 665                 670
```

```
Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr
        675                 680                 685

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
    690                 695                 700

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
705                 710                 715                 720

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
                725                 730                 735

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
            740                 745                 750

Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
        755                 760                 765

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
    770                 775                 780

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
785                 790                 795                 800

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
                805                 810                 815

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr
            820                 825                 830

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
        835                 840                 845

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
    850                 855                 860

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
865                 870                 875                 880

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu
                885                 890                 895

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
            900                 905                 910

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
        915                 920                 925

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
    930                 935                 940
```

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
945 950 955 960

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
965 970 975

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
980 985 990

Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
995 1000 1005

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu
1010 1015 1020

Pro

<210> 103
<211> 1024
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 103
Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
1 5 10 15

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
20 25 30

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
35 40 45

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr
50 55 60

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
65 70 75 80

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
85 90 95

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
100 105 110

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
115 120 125

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
130 135 140

```
Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
145                 150                 155                 160

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
                165                 170                 175

Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
            180                 185                 190

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
        195                 200                 205

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
    210                 215                 220

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
225                 230                 235                 240

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
                245                 250                 255

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
            260                 265                 270

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
        275                 280                 285

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
    290                 295                 300

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
305                 310                 315                 320

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
                325                 330                 335

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
            340                 345                 350

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu
        355                 360                 365

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
    370                 375                 380

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
385                 390                 395                 400

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
                405                 410                 415
```

```
Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
            420                 425                 430

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr
        435                 440                 445

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
    450                 455                 460

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
465                 470                 475                 480

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
                485                 490                 495

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
            500                 505                 510

Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
        515                 520                 525

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
    530                 535                 540

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
545                 550                 555                 560

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
                565                 570                 575

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
            580                 585                 590

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
        595                 600                 605

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
    610                 615                 620

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
625                 630                 635                 640

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
                645                 650                 655

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            660                 665                 670

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
        675                 680                 685
```

```
Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
    690                 695                 700

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
705                 710                 715                 720

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
                725                 730                 735

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
            740                 745                 750

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
        755                 760                 765

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu
    770                 775                 780

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
785                 790                 795                 800

Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
                805                 810                 815

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr
            820                 825                 830

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
        835                 840                 845

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
    850                 855                 860

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
865                 870                 875                 880

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
                885                 890                 895

Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
            900                 905                 910

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
        915                 920                 925

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
    930                 935                 940

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
945                 950                 955                 960
```

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
965 970 975

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
980 985 990

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
995 1000 1005

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
1010 1015 1020

Thr

<210> 104
<211> 1024
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 104
Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
1 5 10 15

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
35 40 45

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
50 55 60

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
65 70 75 80

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
85 90 95

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
100 105 110

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
115 120 125

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
130 135 140

```
Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
145                 150                 155                 160

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
                165                 170                 175

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
            180                 185                 190

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
        195                 200                 205

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
    210                 215                 220

Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
225                 230                 235                 240

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
                245                 250                 255

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
            260                 265                 270

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        275                 280                 285

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
    290                 295                 300

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
305                 310                 315                 320

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
                325                 330                 335

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
            340                 345                 350

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
        355                 360                 365

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
    370                 375                 380

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
385                 390                 395                 400

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
                405                 410                 415
```

```
Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
            420                 425                 430

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
        435                 440                 445

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu
    450                 455                 460

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
465                 470                 475                 480

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
                485                 490                 495

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
            500                 505                 510

Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
        515                 520                 525

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
    530                 535                 540

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
545                 550                 555                 560

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
                565                 570                 575

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu
            580                 585                 590

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
        595                 600                 605

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
    610                 615                 620

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
625                 630                 635                 640

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
                645                 650                 655

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            660                 665                 670

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
        675                 680                 685
```

```
Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
    690                 695                 700

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
705                 710                 715                 720

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
                725                 730                 735

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu
            740                 745                 750

Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
        755                 760                 765

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
    770                 775                 780

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
785                 790                 795                 800

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
                805                 810                 815

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
            820                 825                 830

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
        835                 840                 845

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
    850                 855                 860

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
865                 870                 875                 880

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
                885                 890                 895

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
            900                 905                 910

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
        915                 920                 925

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
    930                 935                 940

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
945                 950                 955                 960
```

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
965 970 975

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
980 985 990

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
995 1000 1005

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro
1010 1015 1020

Ala

<210> 105
<211> 1024
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 105
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
1 5 10 15

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
35 40 45

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala
50 55 60

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
65 70 75 80

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
85 90 95

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
100 105 110

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
115 120 125

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
130 135 140

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
145 150 155 160

```
Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
                165                 170                 175

Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
            180                 185                 190

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
        195                 200                 205

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
    210                 215                 220

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
225                 230                 235                 240

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala
                245                 250                 255

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
            260                 265                 270

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
        275                 280                 285

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu
    290                 295                 300

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
305                 310                 315                 320

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
                325                 330                 335

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr
            340                 345                 350

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
        355                 360                 365

Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
    370                 375                 380

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
385                 390                 395                 400

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
                405                 410                 415

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
            420                 425                 430
```

```
Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
        435                 440                 445

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
    450                 455                 460

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
465                 470                 475                 480

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
                485                 490                 495

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
            500                 505                 510

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        515                 520                 525

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr
    530                 535                 540

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
545                 550                 555                 560

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
                565                 570                 575

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
            580                 585                 590

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
        595                 600                 605

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
    610                 615                 620

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
625                 630                 635                 640

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
                645                 650                 655

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
            660                 665                 670

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
        675                 680                 685

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
    690                 695                 700
```

```
Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
705                 710                 715                 720

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
                725                 730                 735

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
            740                 745                 750

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
        755                 760                 765

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
    770                 775                 780

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu
785                 790                 795                 800

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
                805                 810                 815

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala
            820                 825                 830

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
        835                 840                 845

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
    850                 855                 860

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
865                 870                 875                 880

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
                885                 890                 895

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
            900                 905                 910

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
        915                 920                 925

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
    930                 935                 940

Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
945                 950                 955                 960

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
                965                 970                 975
```

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
980 985 990

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
995 1000 1005

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro
1010 1015 1020

Ala

<210> 106
<211> 1024
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 106
Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr
1 5 10 15

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
20 25 30

Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
35 40 45

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu
50 55 60

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
65 70 75 80

Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
85 90 95

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
100 105 110

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
115 120 125

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
130 135 140

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
145 150 155 160

```
Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
                165                 170                 175

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
            180                 185                 190

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
        195                 200                 205

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
    210                 215                 220

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
225                 230                 235                 240

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
                245                 250                 255

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
            260                 265                 270

Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
        275                 280                 285

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
    290                 295                 300

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
305                 310                 315                 320

Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
                325                 330                 335

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu
            340                 345                 350

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
        355                 360                 365

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    370                 375                 380

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
385                 390                 395                 400

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
                405                 410                 415

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
            420                 425                 430
```

```
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
        435                 440                 445

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
    450                 455                 460

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
465                 470                 475                 480

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
                485                 490                 495

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu
            500                 505                 510

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        515                 520                 525

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala
    530                 535                 540

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
545                 550                 555                 560

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
                565                 570                 575

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
            580                 585                 590

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
        595                 600                 605

Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    610                 615                 620

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
625                 630                 635                 640

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
                645                 650                 655

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
            660                 665                 670

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
        675                 680                 685

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
    690                 695                 700
```

```
Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
705                 710                 715                 720

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu
                725                 730                 735

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
            740                 745                 750

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
        755                 760                 765

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
    770                 775                 780

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
785                 790                 795                 800

Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
                805                 810                 815

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
            820                 825                 830

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
        835                 840                 845

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    850                 855                 860

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
865                 870                 875                 880

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
                885                 890                 895

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
            900                 905                 910

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
        915                 920                 925

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
    930                 935                 940

Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
945                 950                 955                 960

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
                965                 970                 975
```

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
980 985 990

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
995 1000 1005

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
1010 1015 1020

Thr

<210> 107
<211> 1024
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 107
Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
1 5 10 15

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
20 25 30

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
35 40 45

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
50 55 60

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
65 70 75 80

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
85 90 95

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
100 105 110

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
115 120 125

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
130 135 140

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
145 150 155 160

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
165 170 175

```
Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
            180                 185                 190

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
        195                 200                 205

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
    210                 215                 220

Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
225                 230                 235                 240

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
                245                 250                 255

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
            260                 265                 270

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        275                 280                 285

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
    290                 295                 300

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
305                 310                 315                 320

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
                325                 330                 335

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
            340                 345                 350

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
        355                 360                 365

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    370                 375                 380

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
385                 390                 395                 400

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
                405                 410                 415

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            420                 425                 430

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
        435                 440                 445
```

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
450 455 460

Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
465 470 475 480

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
485 490 495

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
500 505 510

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
515 520 525

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
530 535 540

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
545 550 555 560

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
565 570 575

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
580 585 590

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
595 600 605

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
610 615 620

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
625 630 635 640

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
645 650 655

Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
660 665 670

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
675 680 685

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
690 695 700

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
705 710 715 720

```
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
                725                 730                 735

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
            740                 745                 750

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        755                 760                 765

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
    770                 775                 780

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
785                 790                 795                 800

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
                805                 810                 815

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
            820                 825                 830

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
        835                 840                 845

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
    850                 855                 860

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
865                 870                 875                 880

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
                885                 890                 895

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            900                 905                 910

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
        915                 920                 925

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
    930                 935                 940

Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
945                 950                 955                 960

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
                965                 970                 975

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
            980                 985                 990
```

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
995 1000 1005

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
1010 1015 1020

Glu

<210> 108
<211> 1024
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 108
Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
1 5 10 15

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
20 25 30

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
35 40 45

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr
50 55 60

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
65 70 75 80

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
85 90 95

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
100 105 110

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
115 120 125

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
130 135 140

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala
145 150 155 160

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
165 170 175

```
Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
            180                 185                 190

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
        195                 200                 205

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
    210                 215                 220

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
225                 230                 235                 240

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
                245                 250                 255

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
            260                 265                 270

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
        275                 280                 285

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
    290                 295                 300

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
305                 310                 315                 320

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
                325                 330                 335

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr
            340                 345                 350

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
        355                 360                 365

Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
    370                 375                 380

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu
385                 390                 395                 400

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
                405                 410                 415

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
            420                 425                 430

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
        435                 440                 445
```

```
Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
    450                 455                 460

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
465                 470                 475                 480

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr
                485                 490                 495

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
            500                 505                 510

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
        515                 520                 525

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala
    530                 535                 540

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
545                 550                 555                 560

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
                565                 570                 575

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu
            580                 585                 590

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
        595                 600                 605

Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    610                 615                 620

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
625                 630                 635                 640

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
                645                 650                 655

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
            660                 665                 670

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
        675                 680                 685

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu
    690                 695                 700

Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
705                 710                 715                 720
```

```
Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
                725                 730                 735

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
            740                 745                 750

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        755                 760                 765

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
    770                 775                 780

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
785                 790                 795                 800

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
                805                 810                 815

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
            820                 825                 830

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
        835                 840                 845

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
    850                 855                 860

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
865                 870                 875                 880

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
                885                 890                 895

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            900                 905                 910

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
        915                 920                 925

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
    930                 935                 940

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
945                 950                 955                 960

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
                965                 970                 975

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
            980                 985                 990
```

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
995 1000 1005

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu
1010 1015 1020

Pro

<210> 109
<211> 1024
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 109
Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr
1 5 10 15

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
20 25 30

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
35 40 45

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
50 55 60

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
65 70 75 80

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
85 90 95

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
100 105 110

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu
115 120 125

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
130 135 140

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
145 150 155 160

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
165 170 175

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
180 185 190

```
Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu
        195                 200                 205

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
    210                 215                 220

Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
225                 230                 235                 240

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu
                245                 250                 255

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
            260                 265                 270

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
        275                 280                 285

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
    290                 295                 300

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
305                 310                 315                 320

Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
                325                 330                 335

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
            340                 345                 350

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
        355                 360                 365

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
    370                 375                 380

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
385                 390                 395                 400

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
                405                 410                 415

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
            420                 425                 430

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
        435                 440                 445

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
    450                 455                 460
```

```
Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
465                 470                 475                 480

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
                485                 490                 495

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
            500                 505                 510

Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
        515                 520                 525

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu
    530                 535                 540

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
545                 550                 555                 560

Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
                565                 570                 575

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
            580                 585                 590

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
        595                 600                 605

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    610                 615                 620

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
625                 630                 635                 640

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
                645                 650                 655

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
            660                 665                 670

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
        675                 680                 685

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
    690                 695                 700

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
705                 710                 715                 720

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
                725                 730                 735
```

```
Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
            740                 745                 750

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        755                 760                 765

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
    770                 775                 780

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
785                 790                 795                 800

Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
                805                 810                 815

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
            820                 825                 830

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
        835                 840                 845

Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    850                 855                 860

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
865                 870                 875                 880

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
                885                 890                 895

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
            900                 905                 910

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr
        915                 920                 925

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
    930                 935                 940

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
945                 950                 955                 960

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala
                965                 970                 975

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
            980                 985                 990

Glu Ser Gly Pro Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
        995                 1000                1005
```

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
1010 1015 1020

Pro

<210> 110
<211> 1024
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 110
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
1 5 10 15

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
35 40 45

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
50 55 60

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
65 70 75 80

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
85 90 95

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr
100 105 110

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
115 120 125

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
130 135 140

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu
145 150 155 160

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
165 170 175

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
180 185 190

```
Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
        195                 200                 205

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
    210                 215                 220

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
225                 230                 235                 240

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr
                245                 250                 255

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
            260                 265                 270

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
        275                 280                 285

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
    290                 295                 300

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
305                 310                 315                 320

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
                325                 330                 335

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
            340                 345                 350

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
        355                 360                 365

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
    370                 375                 380

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
385                 390                 395                 400

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
                405                 410                 415

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
            420                 425                 430

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
        435                 440                 445

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
    450                 455                 460
```

```
Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
465                 470                 475                 480

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu
                485                 490                 495

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
            500                 505                 510

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
        515                 520                 525

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
    530                 535                 540

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu
545                 550                 555                 560

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
                565                 570                 575

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
            580                 585                 590

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
        595                 600                 605

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
    610                 615                 620

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
625                 630                 635                 640

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
                645                 650                 655

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            660                 665                 670

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
        675                 680                 685

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu
    690                 695                 700

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
705                 710                 715                 720

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala
                725                 730                 735
```

```
Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
            740                 745                 750

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
        755                 760                 765

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
    770                 775                 780

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
785                 790                 795                 800

Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
                805                 810                 815

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
            820                 825                 830

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
        835                 840                 845

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
    850                 855                 860

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
865                 870                 875                 880

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
                885                 890                 895

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
            900                 905                 910

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
        915                 920                 925

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
    930                 935                 940

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
945                 950                 955                 960

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
                965                 970                 975

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
            980                 985                 990

Ser Thr Glu Glu Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
        995                 1000                1005
```

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
1010 1015 1020

Glu

<210> 111
<211> 948
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 111
Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
1 5 10 15

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
20 25 30

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
35 40 45

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly
50 55 60

Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
65 70 75 80

Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
85 90 95

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro
100 105 110

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala
115 120 125

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
130 135 140

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr
145 150 155 160

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
165 170 175

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
180 185 190

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
195 200 205

```
Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
    210                 215                 220

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
225                 230                 235                 240

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
                245                 250                 255

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
            260                 265                 270

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
        275                 280                 285

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
    290                 295                 300

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
305                 310                 315                 320

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
                325                 330                 335

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
            340                 345                 350

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
        355                 360                 365

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    370                 375                 380

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
385                 390                 395                 400

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
                405                 410                 415

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
            420                 425                 430

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
        435                 440                 445

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
    450                 455                 460

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
465                 470                 475                 480
```

```
Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
                485                 490                 495

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala
            500                 505                 510

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
        515                 520                 525

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro
    530                 535                 540

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala
545                 550                 555                 560

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                565                 570                 575

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
            580                 585                 590

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
        595                 600                 605

Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
    610                 615                 620

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
625                 630                 635                 640

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
                645                 650                 655

Ser Ser Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
            660                 665                 670

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
        675                 680                 685

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
    690                 695                 700

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
705                 710                 715                 720

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
                725                 730                 735

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
            740                 745                 750
```

```
Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
        755                 760                 765

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
    770                 775                 780

Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala
785                 790                 795                 800

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
                805                 810                 815

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
            820                 825                 830

Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
        835                 840                 845

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
    850                 855                 860

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
865                 870                 875                 880

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
                885                 890                 895

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
            900                 905                 910

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
        915                 920                 925

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
    930                 935                 940

Thr Gly Ser Pro
945
```

<210> 112
<211> 1024
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 112

```
Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly
1               5                   10                  15
```

```
Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
            20                  25                  30

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
        35                  40                  45

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
    50                  55                  60

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
65                  70                  75                  80

Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
                85                  90                  95

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
            100                 105                 110

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
        115                 120                 125

Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
    130                 135                 140

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
145                 150                 155                 160

Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala
                165                 170                 175

Ser Ser Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
            180                 185                 190

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
        195                 200                 205

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
    210                 215                 220

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
225                 230                 235                 240

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
                245                 250                 255

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
            260                 265                 270

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
        275                 280                 285
```

```
Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
    290                 295                 300

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
305                 310                 315                 320

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
                325                 330                 335

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro
            340                 345                 350

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala
        355                 360                 365

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    370                 375                 380

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
385                 390                 395                 400

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
                405                 410                 415

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
            420                 425                 430

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
        435                 440                 445

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
    450                 455                 460

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
465                 470                 475                 480

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
                485                 490                 495

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
            500                 505                 510

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
        515                 520                 525

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro
    530                 535                 540

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
545                 550                 555                 560
```

```
Ser Ser Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
                565                 570                 575

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
            580                 585                 590

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
        595                 600                 605

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
    610                 615                 620

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro
625                 630                 635                 640

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
                645                 650                 655

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            660                 665                 670

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
        675                 680                 685

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
    690                 695                 700

Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
705                 710                 715                 720

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
                725                 730                 735

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
            740                 745                 750

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
        755                 760                 765

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
    770                 775                 780

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
785                 790                 795                 800

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
                805                 810                 815

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
            820                 825                 830
```

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
835 840 845

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
850 855 860

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
865 870 875 880

Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
885 890 895

Ser Ser Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
900 905 910

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
915 920 925

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
930 935 940

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
945 950 955 960

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro
965 970 975

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
980 985 990

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
995 1000 1005

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser
1010 1015 1020

Thr

<210> 113
<211> 1024
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 113
Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
1 5 10 15

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala
20 25 30

```
Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
        35                  40                  45

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
    50                  55                  60

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
65                  70                  75                  80

Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                85                  90                  95

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly
            100                 105                 110

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
        115                 120                 125

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
    130                 135                 140

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
145                 150                 155                 160

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
                165                 170                 175

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            180                 185                 190

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
        195                 200                 205

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
    210                 215                 220

Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
225                 230                 235                 240

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
                245                 250                 255

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
            260                 265                 270

Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
        275                 280                 285

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr
    290                 295                 300
```

```
Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
305                 310                 315                 320

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
                325                 330                 335

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
            340                 345                 350

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
        355                 360                 365

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    370                 375                 380

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
385                 390                 395                 400

Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
                405                 410                 415

Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
            420                 425                 430

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly
        435                 440                 445

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
    450                 455                 460

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
465                 470                 475                 480

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro
                485                 490                 495

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
            500                 505                 510

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
        515                 520                 525

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
    530                 535                 540

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
545                 550                 555                 560

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                565                 570                 575
```

```
Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
            580                 585                 590

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
        595                 600                 605

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
    610                 615                 620

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
625                 630                 635                 640

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
                645                 650                 655

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
            660                 665                 670

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
        675                 680                 685

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
    690                 695                 700

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
705                 710                 715                 720

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
                725                 730                 735

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
            740                 745                 750

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
        755                 760                 765

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
    770                 775                 780

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
785                 790                 795                 800

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
                805                 810                 815

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
            820                 825                 830

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala
        835                 840                 845
```

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
850 855 860

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
865 870 875 880

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala
885 890 895

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
900 905 910

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
915 920 925

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
930 935 940

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
945 950 955 960

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
965 970 975

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
980 985 990

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
995 1000 1005

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
1010 1015 1020

Thr

<210> 114
<211> 1024
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 114
Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
1 5 10 15

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
20 25 30

```
Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
        35                  40                  45

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
    50                  55                  60

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
65                  70                  75                  80

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
                85                  90                  95

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
            100                 105                 110

Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
        115                 120                 125

Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
    130                 135                 140

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr
145                 150                 155                 160

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
                165                 170                 175

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            180                 185                 190

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly
        195                 200                 205

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
    210                 215                 220

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
225                 230                 235                 240

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
                245                 250                 255

Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
            260                 265                 270

Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
        275                 280                 285

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
    290                 295                 300
```

```
Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
305                 310                 315                 320

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
                325                 330                 335

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
            340                 345                 350

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala
        355                 360                 365

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    370                 375                 380

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
385                 390                 395                 400

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
                405                 410                 415

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
            420                 425                 430

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
        435                 440                 445

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
    450                 455                 460

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
465                 470                 475                 480

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
                485                 490                 495

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
            500                 505                 510

Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
        515                 520                 525

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
    530                 535                 540

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
545                 550                 555                 560

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                565                 570                 575
```

```
Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
            580                 585                 590

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
        595                 600                 605

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
    610                 615                 620

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly
625                 630                 635                 640

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
                645                 650                 655

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
            660                 665                 670

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
        675                 680                 685

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
    690                 695                 700

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
705                 710                 715                 720

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
                725                 730                 735

Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala
            740                 745                 750

Ser Ser Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
        755                 760                 765

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
    770                 775                 780

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
785                 790                 795                 800

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
                805                 810                 815

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
            820                 825                 830

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
        835                 840                 845
```

Ser Ser Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
850 855 860

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
865 870 875 880

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
885 890 895

Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
900 905 910

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
915 920 925

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
930 935 940

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
945 950 955 960

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
965 970 975

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
980 985 990

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
995 1000 1005

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
1010 1015 1020

Pro

<210> 115
<211> 1024
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 115
Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
1 5 10 15

Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala
20 25 30

Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
35 40 45

```
Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
    50                  55                  60

Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala
65                  70                  75                  80

Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                85                  90                  95

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
            100                 105                 110

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala
        115                 120                 125

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    130                 135                 140

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
145                 150                 155                 160

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
                165                 170                 175

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
            180                 185                 190

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
        195                 200                 205

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
    210                 215                 220

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
225                 230                 235                 240

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
                245                 250                 255

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
            260                 265                 270

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
        275                 280                 285

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
    290                 295                 300

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
305                 310                 315                 320
```

```
Ser Ser Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
                325                 330                 335

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
            340                 345                 350

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
        355                 360                 365

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    370                 375                 380

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr
385                 390                 395                 400

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
                405                 410                 415

Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
            420                 425                 430

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro
        435                 440                 445

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala
    450                 455                 460

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
465                 470                 475                 480

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
                485                 490                 495

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
            500                 505                 510

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
        515                 520                 525

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
    530                 535                 540

Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
545                 550                 555                 560

Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
                565                 570                 575

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
            580                 585                 590
```

```
Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
        595                 600                 605

Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
    610                 615                 620

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly
625                 630                 635                 640

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
                645                 650                 655

Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
            660                 665                 670

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
        675                 680                 685

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
    690                 695                 700

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
705                 710                 715                 720

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
                725                 730                 735

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
            740                 745                 750

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
        755                 760                 765

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
    770                 775                 780

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
785                 790                 795                 800

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
                805                 810                 815

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro
            820                 825                 830

Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala
        835                 840                 845

Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    850                 855                 860
```

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
865 870 875 880

Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
885 890 895

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
900 905 910

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
915 920 925

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
930 935 940

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
945 950 955 960

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
965 970 975

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
980 985 990

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
995 1000 1005

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
1010 1015 1020

Gly

<210> 116
<211> 1024
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 116
Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
1 5 10 15

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
20 25 30

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
35 40 45

```
Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly
    50                  55                  60

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
65                  70                  75                  80

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
                85                  90                  95

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
            100                 105                 110

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala
        115                 120                 125

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    130                 135                 140

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
145                 150                 155                 160

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
                165                 170                 175

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
            180                 185                 190

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly
        195                 200                 205

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
    210                 215                 220

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
225                 230                 235                 240

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly
                245                 250                 255

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
            260                 265                 270

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
        275                 280                 285

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
    290                 295                 300

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
305                 310                 315                 320
```

```
Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
               325                 330                 335

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
            340                 345                 350

Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
        355                 360                 365

Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
    370                 375                 380

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
385                 390                 395                 400

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
               405                 410                 415

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            420                 425                 430

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
        435                 440                 445

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
    450                 455                 460

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
465                 470                 475                 480

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
               485                 490                 495

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
            500                 505                 510

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
        515                 520                 525

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
    530                 535                 540

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
545                 550                 555                 560

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
               565                 570                 575

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly
            580                 585                 590
```

```
Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
        595                 600                 605

Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
    610                 615                 620

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
625                 630                 635                 640

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
                645                 650                 655

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            660                 665                 670

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
        675                 680                 685

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
    690                 695                 700

Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
705                 710                 715                 720

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
                725                 730                 735

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
            740                 745                 750

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
        755                 760                 765

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
    770                 775                 780

Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala
785                 790                 795                 800

Ser Ser Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
                805                 810                 815

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
            820                 825                 830

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
        835                 840                 845

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
    850                 855                 860
```

```
Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro
865                 870                 875                 880

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala
                885                 890                 895

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
            900                 905                 910

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr
        915                 920                 925

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
    930                 935                 940

Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
945                 950                 955                 960

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
                965                 970                 975

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
            980                 985                 990

Gly Thr Gly Pro Gly Ser Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro
        995                 1000                1005

Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser Ser Pro  Gly Ala Ser
    1010                1015                1020

Pro
```

<210> 117
<211> 1024
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 117

```
Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
1               5                   10                  15

Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
            20                  25                  30

Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
        35                  40                  45

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
    50                  55                  60
```

```
Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
65                  70                  75                  80

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
                85                  90                  95

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
            100                 105                 110

Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
        115                 120                 125

Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
    130                 135                 140

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
145                 150                 155                 160

Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala
                165                 170                 175

Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
            180                 185                 190

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro
        195                 200                 205

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
    210                 215                 220

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
225                 230                 235                 240

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro
                245                 250                 255

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
            260                 265                 270

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
        275                 280                 285

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
    290                 295                 300

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
305                 310                 315                 320

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
                325                 330                 335
```

```
Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
            340                 345                 350

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
        355                 360                 365

Ser Ser Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
    370                 375                 380

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
385                 390                 395                 400

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
                405                 410                 415

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
            420                 425                 430

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
        435                 440                 445

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
    450                 455                 460

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
465                 470                 475                 480

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
                485                 490                 495

Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala
            500                 505                 510

Ser Ser Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
        515                 520                 525

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
    530                 535                 540

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
545                 550                 555                 560

Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                565                 570                 575

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly
            580                 585                 590

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
        595                 600                 605
```

```
Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
    610                 615                 620

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro
625                 630                 635                 640

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala
                645                 650                 655

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
            660                 665                 670

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
        675                 680                 685

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
    690                 695                 700

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
705                 710                 715                 720

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
                725                 730                 735

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
            740                 745                 750

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
        755                 760                 765

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly
    770                 775                 780

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
785                 790                 795                 800

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
                805                 810                 815

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
            820                 825                 830

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
        835                 840                 845

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
    850                 855                 860

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro
865                 870                 875                 880
```

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
885 890 895

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
900 905 910

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
915 920 925

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
930 935 940

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
945 950 955 960

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
965 970 975

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
980 985 990

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
995 1000 1005

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro
1010 1015 1020

Gly

<210> 118
<211> 1024
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 118
Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
1 5 10 15

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
20 25 30

Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
35 40 45

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
50 55 60

```
Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
65                  70                  75                  80

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                85                  90                  95

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
            100                 105                 110

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
        115                 120                 125

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
    130                 135                 140

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
145                 150                 155                 160

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
                165                 170                 175

Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
            180                 185                 190

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
        195                 200                 205

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
    210                 215                 220

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
225                 230                 235                 240

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
                245                 250                 255

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
            260                 265                 270

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
        275                 280                 285

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
    290                 295                 300

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala
305                 310                 315                 320

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
                325                 330                 335
```

```
Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
            340                 345                 350

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
        355                 360                 365

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
    370                 375                 380

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
385                 390                 395                 400

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
                405                 410                 415

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            420                 425                 430

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro
        435                 440                 445

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala
    450                 455                 460

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
465                 470                 475                 480

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
                485                 490                 495

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
            500                 505                 510

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
        515                 520                 525

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro
    530                 535                 540

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
545                 550                 555                 560

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
                565                 570                 575

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
            580                 585                 590

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
        595                 600                 605
```

```
Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
    610                 615                 620

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr
625                 630                 635                 640

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
                645                 650                 655

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
            660                 665                 670

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
        675                 680                 685

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
    690                 695                 700

Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
705                 710                 715                 720

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
                725                 730                 735

Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala
            740                 745                 750

Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
        755                 760                 765

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr
    770                 775                 780

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
785                 790                 795                 800

Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
                805                 810                 815

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
            820                 825                 830

Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
        835                 840                 845

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
    850                 855                 860

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
865                 870                 875                 880
```

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
885 890 895

Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
900 905 910

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
915 920 925

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala
930 935 940

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
945 950 955 960

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
965 970 975

Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
980 985 990

Ser Ser Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
995 1000 1005

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
1010 1015 1020

Thr

<210> 119
<211> 1024
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 119
Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
1 5 10 15

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala
20 25 30

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
35 40 45

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
50 55 60

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
65 70 75 80

```
Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
                85                  90                  95

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
            100                 105                 110

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
        115                 120                 125

Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
    130                 135                 140

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
145                 150                 155                 160

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
                165                 170                 175

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
            180                 185                 190

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
        195                 200                 205

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
    210                 215                 220

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
225                 230                 235                 240

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
                245                 250                 255

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
            260                 265                 270

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
        275                 280                 285

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
    290                 295                 300

Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
305                 310                 315                 320

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
                325                 330                 335

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro
            340                 345                 350
```

```
Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
        355                 360                 365

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    370                 375                 380

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro
385                 390                 395                 400

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
                405                 410                 415

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
            420                 425                 430

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
        435                 440                 445

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
    450                 455                 460

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
465                 470                 475                 480

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
                485                 490                 495

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
            500                 505                 510

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
        515                 520                 525

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
    530                 535                 540

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
545                 550                 555                 560

Ser Ser Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
                565                 570                 575

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
            580                 585                 590

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
        595                 600                 605

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
    610                 615                 620
```

```
Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
625                 630                 635                 640

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
                645                 650                 655

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            660                 665                 670

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
        675                 680                 685

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
    690                 695                 700

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
705                 710                 715                 720

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
                725                 730                 735

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
            740                 745                 750

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
        755                 760                 765

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
    770                 775                 780

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
785                 790                 795                 800

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
                805                 810                 815

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr
            820                 825                 830

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
        835                 840                 845

Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
    850                 855                 860

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
865                 870                 875                 880

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
                885                 890                 895
```

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
900 905 910

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
915 920 925

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
930 935 940

Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
945 950 955 960

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
965 970 975

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
980 985 990

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
995 1000 1005

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
1010 1015 1020

Gly

<210> 120
<211> 1024
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 120
Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
1 5 10 15

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
20 25 30

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
35 40 45

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro
50 55 60

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
65 70 75 80

```
Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
                85                  90                  95

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro
            100                 105                 110

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
        115                 120                 125

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
    130                 135                 140

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
145                 150                 155                 160

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
                165                 170                 175

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
            180                 185                 190

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
        195                 200                 205

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
    210                 215                 220

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
225                 230                 235                 240

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro
                245                 250                 255

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala
            260                 265                 270

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
        275                 280                 285

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
    290                 295                 300

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
305                 310                 315                 320

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
                325                 330                 335

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
            340                 345                 350
```

```
Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
        355                 360                 365

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
    370                 375                 380

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
385                 390                 395                 400

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
                405                 410                 415

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
            420                 425                 430

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro
        435                 440                 445

Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala
    450                 455                 460

Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
465                 470                 475                 480

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
                485                 490                 495

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
            500                 505                 510

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
        515                 520                 525

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
    530                 535                 540

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
545                 550                 555                 560

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                565                 570                 575

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
            580                 585                 590

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
        595                 600                 605

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
    610                 615                 620
```

```
Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
625                 630                 635                 640

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
                645                 650                 655

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
            660                 665                 670

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
        675                 680                 685

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
    690                 695                 700

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
705                 710                 715                 720

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
                725                 730                 735

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
            740                 745                 750

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
        755                 760                 765

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
    770                 775                 780

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
785                 790                 795                 800

Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
                805                 810                 815

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
            820                 825                 830

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
        835                 840                 845

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
    850                 855                 860

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
865                 870                 875                 880

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
                885                 890                 895
```

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
900 905 910

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
915 920 925

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
930 935 940

Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
945 950 955 960

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
965 970 975

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
980 985 990

Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
995 1000 1005

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser
1010 1015 1020

Pro

<210> 121
<211> 1024
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 121
Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
1 5 10 15

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
20 25 30

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
35 40 45

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
50 55 60

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
65 70 75 80

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
85 90 95

```
Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly
            100                 105                 110

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
        115                 120                 125

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
    130                 135                 140

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro
145                 150                 155                 160

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala
                165                 170                 175

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            180                 185                 190

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
        195                 200                 205

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
    210                 215                 220

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
225                 230                 235                 240

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly
                245                 250                 255

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
            260                 265                 270

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
        275                 280                 285

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro
    290                 295                 300

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala
305                 310                 315                 320

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                325                 330                 335

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
            340                 345                 350

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
        355                 360                 365
```

```
Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
    370                 375                 380

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
385                 390                 395                 400

Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala
                405                 410                 415

Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
            420                 425                 430

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
        435                 440                 445

Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
    450                 455                 460

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
465                 470                 475                 480

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
                485                 490                 495

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
            500                 505                 510

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
        515                 520                 525

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
    530                 535                 540

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala
545                 550                 555                 560

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                565                 570                 575

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
            580                 585                 590

Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
        595                 600                 605

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
    610                 615                 620

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
625                 630                 635                 640
```

```
Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
                645                 650                 655

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
            660                 665                 670

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro
        675                 680                 685

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
    690                 695                 700

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
705                 710                 715                 720

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly
                725                 730                 735

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
            740                 745                 750

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
        755                 760                 765

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
    770                 775                 780

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala
785                 790                 795                 800

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
                805                 810                 815

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
            820                 825                 830

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
        835                 840                 845

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
    850                 855                 860

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro
865                 870                 875                 880

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
                885                 890                 895

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
            900                 905                 910
```

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr
915 920 925

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
930 935 940

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
945 950 955 960

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
965 970 975

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
980 985 990

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
995 1000 1005

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser
1010 1015 1020

Pro

<210> 122
<211> 1024
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 122
Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
1 5 10 15

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
20 25 30

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
35 40 45

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
50 55 60

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
65 70 75 80

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
85 90 95

```
Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
            100                 105                 110

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
        115                 120                 125

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
    130                 135                 140

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly
145                 150                 155                 160

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
                165                 170                 175

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
            180                 185                 190

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
        195                 200                 205

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
    210                 215                 220

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
225                 230                 235                 240

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
                245                 250                 255

Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
            260                 265                 270

Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
        275                 280                 285

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro
    290                 295                 300

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
305                 310                 315                 320

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
                325                 330                 335

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
            340                 345                 350

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
        355                 360                 365
```

```
Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
    370                 375                 380

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
385                 390                 395                 400

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
                405                 410                 415

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
            420                 425                 430

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
        435                 440                 445

Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
    450                 455                 460

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
465                 470                 475                 480

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
                485                 490                 495

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
            500                 505                 510

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
        515                 520                 525

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr
    530                 535                 540

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
545                 550                 555                 560

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                565                 570                 575

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
            580                 585                 590

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
        595                 600                 605

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
    610                 615                 620

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro
625                 630                 635                 640
```

```
Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala
               645                 650                 655

Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
            660                 665                 670

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
        675                 680                 685

Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
    690                 695                 700

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
705                 710                 715                 720

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
               725                 730                 735

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
            740                 745                 750

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
        755                 760                 765

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
    770                 775                 780

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala
785                 790                 795                 800

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
               805                 810                 815

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
            820                 825                 830

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
        835                 840                 845

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    850                 855                 860

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly
865                 870                 875                 880

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
               885                 890                 895

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
            900                 905                 910
```

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
915 920 925

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
930 935 940

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
945 950 955 960

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
965 970 975

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
980 985 990

Ser Ser Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
995 1000 1005

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
1010 1015 1020

Pro

<210> 123
<211> 72
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 123
Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
1 5 10 15

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
20 25 30

Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
35 40 45

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
50 55 60

Thr Ser Gly Ser Glu Thr Pro Gly
65 70

<210> 124
<211> 72
<212> PRT
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 124
Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser
1 5 10 15

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
20 25 30

Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
35 40 45

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu
50 55 60

Pro Ser Glu Gly Ser Ala Pro Gly
65 70

<210> 125
<211> 108
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 125
Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
1 5 10 15

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
20 25 30

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
35 40 45

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
50 55 60

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
65 70 75 80

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
85 90 95

Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
100 105

<210> 126
<211> 108
<212> PRT
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 126
Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
1 5 10 15

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
20 25 30

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
35 40 45

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
50 55 60

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
65 70 75 80

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
85 90 95

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
100 105

<210> 127
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 127
Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser
1 5 10 15

Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
20 25 30

Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
35 40 45

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala
50 55 60

Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu
65 70 75 80

Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
85 90 95

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala
100 105 110

Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr
115 120 125

Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser
130 135 140

<210> 128
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 128
Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
1 5 10 15

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
20 25 30

Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
35 40 45

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
50 55 60

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
65 70 75 80

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
85 90 95

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly
100 105 110

Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser
115 120 125

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
130 135 140

<210> 129
<211> 180
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 129
Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1 5 10 15

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser
20 25 30

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro
35 40 45

Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
50 55 60

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
65 70 75 80

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
85 90 95

Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
100 105 110

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu
115 120 125

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro
130 135 140

Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
145 150 155 160

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu
165 170 175

Pro Ala Thr Ser
180

<210> 130
<211> 216
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 130
Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
1 5 10 15

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
20 25 30

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
35 40 45

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
50 55 60

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro
65 70 75 80

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
85 90 95

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
100 105 110

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
115 120 125

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
130 135 140

Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
145 150 155 160

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
165 170 175

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
180 185 190

Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
195 200 205

Pro Gly Thr Ser Glu Ser Ala Thr
210 215

<210> 131
<211> 252
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 131
Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
1 5 10 15

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
20 25 30

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
35 40 45

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
50 55 60

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
65 70 75 80

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
85 90 95

Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
100 105 110

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
115 120 125

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
130 135 140

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
145 150 155 160

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
165 170 175

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
180 185 190

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
195 200 205

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
210 215 220

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
225 230 235 240

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
245 250

<210> 132
<211> 288
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 132
Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
1 5 10 15

Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
20 25 30

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala
35 40 45

Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu
50 55 60

Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
65 70 75 80

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser
85 90 95

Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
100 105 110

Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr
115 120 125

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser
130 135 140

Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr
145 150 155 160

Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
165 170 175

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala
180 185 190

Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
195 200 205

Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser
210 215 220

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser
225 230 235 240

Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
245 250 255

Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
260 265 270

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala
275 280 285

<210> 133
<211> 324
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 133

```
Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
1               5                   10                  15

Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro
            20                  25                  30

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr
        35                  40                  45

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
    50                  55                  60

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
65                  70                  75                  80

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
                85                  90                  95

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
            100                 105                 110

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
        115                 120                 125

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro
    130                 135                 140

Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
145                 150                 155                 160

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser
                165                 170                 175

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro
            180                 185                 190

Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
        195                 200                 205

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
    210                 215                 220

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
225                 230                 235                 240

Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
                245                 250                 255
```

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu
260 265 270

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro
275 280 285

Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
290 295 300

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu
305 310 315 320

Pro Ala Thr Ser

<210> 134
<211> 360
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 134
Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
1 5 10 15

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro
20 25 30

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro
35 40 45

Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
50 55 60

Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
65 70 75 80

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
85 90 95

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
100 105 110

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
115 120 125

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
130 135 140

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
145 150 155 160

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
165 170 175

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
180 185 190

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
195 200 205

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro
210 215 220

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
225 230 235 240

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
245 250 255

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
260 265 270

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
275 280 285

Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
290 295 300

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
305 310 315 320

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
325 330 335

Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
340 345 350

Pro Gly Thr Ser Glu Ser Ala Thr
355 360

<210> 135
<211> 360
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 135
Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1 5 10 15

```
Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
            20                  25                  30

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
        35                  40                  45

Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
    50                  55                  60

Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro
65                  70                  75                  80

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr
                85                  90                  95

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            100                 105                 110

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
        115                 120                 125

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
    130                 135                 140

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
145                 150                 155                 160

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
                165                 170                 175

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro
            180                 185                 190

Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
        195                 200                 205

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser
    210                 215                 220

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro
225                 230                 235                 240

Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
                245                 250                 255

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
            260                 265                 270

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
        275                 280                 285
```

Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
290 295 300

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu
305 310 315 320

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro
325 330 335

Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
340 345 350

Pro Gly Thr Ser Thr Glu Pro Ser
355 360

<210> 136
<211> 432
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 136
Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
1 5 10 15

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
20 25 30

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
35 40 45

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
50 55 60

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser
65 70 75 80

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser
85 90 95

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
100 105 110

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro
115 120 125

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro
130 135 140

```
Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
145                 150                 155                 160

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
                165                 170                 175

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
            180                 185                 190

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
        195                 200                 205

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser
    210                 215                 220

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser
225                 230                 235                 240

Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
                245                 250                 255

Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
            260                 265                 270

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
        275                 280                 285

Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
    290                 295                 300

Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
305                 310                 315                 320

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
                325                 330                 335

Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
            340                 345                 350

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
        355                 360                 365

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser
    370                 375                 380

Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
385                 390                 395                 400

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
                405                 410                 415
```

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser
420 425 430

<210> 137
<211> 468
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 137
Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
1 5 10 15

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro
20 25 30

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
35 40 45

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
50 55 60

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser
65 70 75 80

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
85 90 95

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
100 105 110

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
115 120 125

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro
130 135 140

Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
145 150 155 160

Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
165 170 175

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser
180 185 190

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
195 200 205

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser
210 215 220

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
225 230 235 240

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
245 250 255

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro
260 265 270

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr
275 280 285

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
290 295 300

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro
305 310 315 320

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro
325 330 335

Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
340 345 350

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
355 360 365

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
370 375 380

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
385 390 395 400

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro
405 410 415

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
420 425 430

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
435 440 445

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser
450 455 460

Glu Ser Ala Thr
465

<210> 138
<211> 504

<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 138

```
Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
1               5                  10                  15

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
            20                  25                  30

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
        35                  40                  45

Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
    50                  55                  60

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
65                  70                  75                  80

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
                85                  90                  95

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
            100                 105                 110

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser
        115                 120                 125

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser
    130                 135                 140

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
145                 150                 155                 160

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro
                165                 170                 175

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro
            180                 185                 190

Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
        195                 200                 205

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
    210                 215                 220

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
225                 230                 235                 240
```

```
Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
                245                 250                 255

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser
            260                 265                 270

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser
        275                 280                 285

Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
    290                 295                 300

Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
305                 310                 315                 320

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
                325                 330                 335

Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
            340                 345                 350

Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
        355                 360                 365

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
    370                 375                 380

Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
385                 390                 395                 400

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
                405                 410                 415

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser
            420                 425                 430

Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
        435                 440                 445

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
    450                 455                 460

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser
465                 470                 475                 480

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
                485                 490                 495

Pro Gly Thr Ser Thr Glu Pro Ser
            500
```

<210> 139
<211> 540
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 139

```
Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr
1               5                   10                  15

Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
            20                  25                  30

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala
        35                  40                  45

Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
    50                  55                  60

Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
65                  70                  75                  80

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro
                85                  90                  95

Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
            100                 105                 110

Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
        115                 120                 125

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro
    130                 135                 140

Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
145                 150                 155                 160

Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr
                165                 170                 175

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr
            180                 185                 190

Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
        195                 200                 205

Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
    210                 215                 220

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser
225                 230                 235                 240
```

```
Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr
                245                 250                 255

Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr
            260                 265                 270

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro
        275                 280                 285

Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
    290                 295                 300

Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr
305                 310                 315                 320

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr
                325                 330                 335

Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
            340                 345                 350

Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
        355                 360                 365

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala
    370                 375                 380

Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu
385                 390                 395                 400

Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
                405                 410                 415

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser
            420                 425                 430

Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
        435                 440                 445

Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
    450                 455                 460

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala
465                 470                 475                 480

Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu
                485                 490                 495

Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser
            500                 505                 510
```

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro
515 520 525

Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro
530 535 540

<210> 140
<211> 576
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 140
Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
1 5 10 15

Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr
20 25 30

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr
35 40 45

Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
50 55 60

Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
65 70 75 80

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro
85 90 95

Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
100 105 110

Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
115 120 125

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser
130 135 140

Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
145 150 155 160

Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
165 170 175

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala
180 185 190

```
Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu
        195                 200                 205

Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
    210                 215                 220

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala
225                 230                 235                 240

Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr
                245                 250                 255

Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser
            260                 265                 270

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala
        275                 280                 285

Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
    290                 295                 300

Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
305                 310                 315                 320

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala
                325                 330                 335

Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu
            340                 345                 350

Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
        355                 360                 365

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser
    370                 375                 380

Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
385                 390                 395                 400

Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr
                405                 410                 415

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser
            420                 425                 430

Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr
        435                 440                 445

Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
    450                 455                 460
```

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala
465 470 475 480

Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
485 490 495

Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser
500 505 510

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser
515 520 525

Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
530 535 540

Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
545 550 555 560

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala
565 570 575

<210> 141
<211> 612
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 141
Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
1 5 10 15

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
20 25 30

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
35 40 45

Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
50 55 60

Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
65 70 75 80

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
85 90 95

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
100 105 110

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
115 120 125

```
Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
    130                 135                 140

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
145                 150                 155                 160

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro
                165                 170                 175

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
            180                 185                 190

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
        195                 200                 205

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser
    210                 215                 220

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
225                 230                 235                 240

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
                245                 250                 255

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
            260                 265                 270

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro
        275                 280                 285

Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
    290                 295                 300

Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
305                 310                 315                 320

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser
                325                 330                 335

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
            340                 345                 350

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser
        355                 360                 365

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
    370                 375                 380

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
385                 390                 395                 400
```

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro
405 410 415

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr
420 425 430

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
435 440 445

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro
450 455 460

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro
465 470 475 480

Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
485 490 495

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
500 505 510

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
515 520 525

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
530 535 540

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro
545 550 555 560

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
565 570 575

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
580 585 590

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser
595 600 605

Glu Ser Ala Thr
610

<210> 142
<211> 648
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 142

```
Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
1               5                   10                  15

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
            20                  25                  30

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser
        35                  40                  45

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
    50                  55                  60

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
65                  70                  75                  80

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro
                85                  90                  95

Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
            100                 105                 110

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser
        115                 120                 125

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser
    130                 135                 140

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
145                 150                 155                 160

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
                165                 170                 175

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
            180                 185                 190

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
        195                 200                 205

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
    210                 215                 220

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
225                 230                 235                 240

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
                245                 250                 255

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
            260                 265                 270
```

```
Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
        275                 280                 285

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
    290                 295                 300

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro
305                 310                 315                 320

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro
                325                 330                 335

Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
            340                 345                 350

Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
        355                 360                 365

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
    370                 375                 380

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
385                 390                 395                 400

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
                405                 410                 415

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
            420                 425                 430

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
        435                 440                 445

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
    450                 455                 460

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
465                 470                 475                 480

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
                485                 490                 495

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro
            500                 505                 510

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
        515                 520                 525

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    530                 535                 540
```

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
545 550 555 560

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
565 570 575

Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
580 585 590

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
595 600 605

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
610 615 620

Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
625 630 635 640

Pro Gly Thr Ser Glu Ser Ala Thr
645

<210> 143
<211> 684
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 143
Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
1 5 10 15

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
20 25 30

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
35 40 45

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
50 55 60

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
65 70 75 80

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser
85 90 95

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
100 105 110

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
115 120 125

```
Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro
    130                 135                 140

Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
145                 150                 155                 160

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser
                165                 170                 175

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser
            180                 185                 190

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
        195                 200                 205

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
    210                 215                 220

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
225                 230                 235                 240

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
                245                 250                 255

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            260                 265                 270

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
        275                 280                 285

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
    290                 295                 300

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
305                 310                 315                 320

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
                325                 330                 335

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            340                 345                 350

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro
        355                 360                 365

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro
    370                 375                 380

Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
385                 390                 395                 400
```

```
Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
                405                 410                 415

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
            420                 425                 430

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
        435                 440                 445

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
    450                 455                 460

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
465                 470                 475                 480

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
                485                 490                 495

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            500                 505                 510

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        515                 520                 525

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    530                 535                 540

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro
545                 550                 555                 560

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
                565                 570                 575

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
            580                 585                 590

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
        595                 600                 605

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
    610                 615                 620

Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
625                 630                 635                 640

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
                645                 650                 655

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
            660                 665                 670
```

Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser
675 680

<210> 144
<211> 720
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 144
Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
1 5 10 15

Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
20 25 30

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
35 40 45

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
50 55 60

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
65 70 75 80

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
85 90 95

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
100 105 110

Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
115 120 125

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
130 135 140

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
145 150 155 160

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
165 170 175

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser
180 185 190

Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
195 200 205

```
Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
    210                 215                 220

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
225                 230                 235                 240

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                245                 250                 255

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            260                 265                 270

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
        275                 280                 285

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    290                 295                 300

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
305                 310                 315                 320

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu
                325                 330                 335

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
            340                 345                 350

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
        355                 360                 365

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
    370                 375                 380

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
385                 390                 395                 400

Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
                405                 410                 415

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly
            420                 425                 430

Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser
        435                 440                 445

Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
    450                 455                 460

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
465                 470                 475                 480
```

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
485 490 495

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
500 505 510

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
515 520 525

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
530 535 540

Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
545 550 555 560

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
565 570 575

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
580 585 590

Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
595 600 605

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
610 615 620

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
625 630 635 640

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
645 650 655

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser
660 665 670

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
675 680 685

Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
690 695 700

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu
705 710 715 720

<210> 145
<211> 756
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 145
Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
1 5 10 15

Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
20 25 30

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
35 40 45

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
50 55 60

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
65 70 75 80

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
85 90 95

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
100 105 110

Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
115 120 125

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
130 135 140

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
145 150 155 160

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
165 170 175

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser
180 185 190

Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
195 200 205

Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
210 215 220

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
225 230 235 240

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
245 250 255

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
260 265 270

```
Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
        275                 280                 285

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    290                 295                 300

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
305                 310                 315                 320

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu
                325                 330                 335

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
            340                 345                 350

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
        355                 360                 365

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
    370                 375                 380

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
385                 390                 395                 400

Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
                405                 410                 415

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly
            420                 425                 430

Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser
        435                 440                 445

Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
    450                 455                 460

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
465                 470                 475                 480

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
                485                 490                 495

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
            500                 505                 510

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
        515                 520                 525

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
    530                 535                 540
```

Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
545 550 555 560

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
565 570 575

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
580 585 590

Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
595 600 605

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
610 615 620

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
625 630 635 640

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
645 650 655

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser
660 665 670

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
675 680 685

Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
690 695 700

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu
705 710 715 720

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
725 730 735

Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
740 745 750

Thr Ser Glu Ser
755

<210> 146
<211> 792
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 146

```
Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1               5                   10                  15

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu
            20                  25                  30

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro
        35                  40                  45

Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    50                  55                  60

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
65                  70                  75                  80

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro
                85                  90                  95

Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            100                 105                 110

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
        115                 120                 125

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser
    130                 135                 140

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
145                 150                 155                 160

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser
                165                 170                 175

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
            180                 185                 190

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
        195                 200                 205

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro
    210                 215                 220

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr
225                 230                 235                 240

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
                245                 250                 255

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
            260                 265                 270
```

```
Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
        275                 280                 285

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
    290                 295                 300

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
305                 310                 315                 320

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                325                 330                 335

Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
            340                 345                 350

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
        355                 360                 365

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
    370                 375                 380

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
385                 390                 395                 400

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser
                405                 410                 415

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser
            420                 425                 430

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
        435                 440                 445

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro
    450                 455                 460

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro
465                 470                 475                 480

Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
                485                 490                 495

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
            500                 505                 510

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        515                 520                 525

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    530                 535                 540
```

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser
545 550 555 560

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser
565 570 575

Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
580 585 590

Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
595 600 605

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
610 615 620

Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
625 630 635 640

Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
645 650 655

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
660 665 670

Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
675 680 685

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
690 695 700

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser
705 710 715 720

Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
725 730 735

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
740 745 750

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser
755 760 765

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
770 775 780

Pro Gly Thr Ser Thr Glu Pro Ser
785 790

<210> 147
<211> 828
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 147

```
Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
1               5                   10                  15

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            20                  25                  30

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
        35                  40                  45

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    50                  55                  60

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu
65                  70                  75                  80

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro
                85                  90                  95

Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
            100                 105                 110

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
        115                 120                 125

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro
    130                 135                 140

Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
145                 150                 155                 160

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
                165                 170                 175

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser
            180                 185                 190

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
        195                 200                 205

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser
    210                 215                 220

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
225                 230                 235                 240

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
                245                 250                 255
```

```
Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro
            260                 265                 270

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr
        275                 280                 285

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
    290                 295                 300

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
305                 310                 315                 320

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                325                 330                 335

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            340                 345                 350

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
        355                 360                 365

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
    370                 375                 380

Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
385                 390                 395                 400

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
                405                 410                 415

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
            420                 425                 430

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
        435                 440                 445

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser
    450                 455                 460

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser
465                 470                 475                 480

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
                485                 490                 495

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro
            500                 505                 510

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro
        515                 520                 525
```

```
Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    530                 535                 540

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
545                 550                 555                 560

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
                565                 570                 575

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
            580                 585                 590

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser
        595                 600                 605

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser
    610                 615                 620

Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
625                 630                 635                 640

Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
                645                 650                 655

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
            660                 665                 670

Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
        675                 680                 685

Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
    690                 695                 700

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
705                 710                 715                 720

Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
                725                 730                 735

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
            740                 745                 750

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser
        755                 760                 765

Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
    770                 775                 780

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
785                 790                 795                 800
```

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser
805 810 815

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
820 825

<210> 148
<211> 72
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 148
Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr
1 5 10 15

Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser
20 25 30

Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr
35 40 45

Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr
50 55 60

Pro Gly Ser Gly Thr Ala Ser Ser
65 70

<210> 149
<211> 72
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 149
Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
1 5 10 15

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
20 25 30

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
35 40 45

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
50 55 60

Ser Gly Thr Ala Ser Ser Ser Pro
65 70

<210> 150
<211> 72
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 150
Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr
1 5 10 15

Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr
20 25 30

Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser
35 40 45

Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser
50 55 60

Gly Ala Thr Gly Ser Pro Gly Ala
65 70

<210> 151
<211> 108
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 151
Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
1 5 10 15

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
20 25 30

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
35 40 45

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
50 55 60

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
65 70 75 80

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
85 90 95

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
100 105

<210> 152
<211> 108
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 152
Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser
1 5 10 15

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
20 25 30

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
35 40 45

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
50 55 60

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
65 70 75 80

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
85 90 95

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
100 105

<210> 153
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 153
Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser
1 5 10 15

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr
20 25 30

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
35 40 45

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
50 55 60

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
65 70 75 80

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
85 90 95

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser
100 105 110

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
115 120 125

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
130 135 140

<210> 154
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 154
Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
1 5 10 15

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
20 25 30

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
35 40 45

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
50 55 60

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
65 70 75 80

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
85 90 95

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
100 105 110

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
115 120 125

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
130 135 140

<210> 155
<211> 180
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 155
Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
1 5 10 15

Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
20 25 30

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
35 40 45

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
50 55 60

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65 70 75 80

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
85 90 95

Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
100 105 110

Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
115 120 125

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser
130 135 140

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145 150 155 160

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
165 170 175

Thr Pro Gly Ser
180

<210> 156
<211> 216
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 156
Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
1 5 10 15

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser
20 25 30

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
35 40 45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
50 55 60

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser
65 70 75 80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
85 90 95

Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
100 105 110

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
115 120 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
130 135 140

Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
145 150 155 160

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser
165 170 175

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
180 185 190

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
195 200 205

Pro Gly Ser Ser Thr Pro Ser Gly
210 215

<210> 157
<211> 252
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 157
Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
1 5 10 15

Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
20 25 30

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
35 40 45

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
50 55 60

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65 70 75 80

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
85 90 95

Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
100 105 110

Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
115 120 125

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser
130 135 140

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145 150 155 160

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
165 170 175

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
180 185 190

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
195 200 205

Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
210 215 220

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
225 230 235 240

Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly
245 250

<210> 158
<211> 288
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 158

```
Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
1               5                   10                  15

Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
            20                  25                  30

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
        35                  40                  45

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
    50                  55                  60

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65                  70                  75                  80

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
                85                  90                  95

Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
            100                 105                 110

Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
        115                 120                 125

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser
    130                 135                 140

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145                 150                 155                 160

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
                165                 170                 175

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
            180                 185                 190

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
        195                 200                 205

Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
    210                 215                 220

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
225                 230                 235                 240

Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
                245                 250                 255

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
            260                 265                 270
```

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser
275 280 285

<210> 159
<211> 324
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 159
Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
1 5 10 15

Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
20 25 30

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
35 40 45

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
50 55 60

Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
65 70 75 80

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser
85 90 95

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
100 105 110

Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
115 120 125

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
130 135 140

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
145 150 155 160

Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
165 170 175

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser
180 185 190

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
195 200 205

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
210 215 220

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly
225 230 235 240

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
245 250 255

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
260 265 270

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly
275 280 285

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
290 295 300

Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
305 310 315 320

Ser Ser Thr Pro

<210> 160
<211> 360
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 160
Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
1 5 10 15

Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
20 25 30

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
35 40 45

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
50 55 60

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65 70 75 80

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
85 90 95

Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
100 105 110

Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
115 120 125

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
130 135 140

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145 150 155 160

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
165 170 175

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly
180 185 190

Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
195 200 205

Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
210 215 220

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser
225 230 235 240

Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
245 250 255

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
260 265 270

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser
275 280 285

Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
290 295 300

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
305 310 315 320

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
325 330 335

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
340 345 350

Gly Ser Pro Gly Ala Ser Pro Gly
355 360

<210> 161
<211> 396
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 161

```
Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
1               5                   10                  15

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala
            20                  25                  30

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
        35                  40                  45

Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
    50                  55                  60

Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
65                  70                  75                  80

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
                85                  90                  95

Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
            100                 105                 110

Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser
        115                 120                 125

Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly
    130                 135                 140

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
145                 150                 155                 160

Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser
                165                 170                 175

Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser
            180                 185                 190

Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr
        195                 200                 205

Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala
    210                 215                 220

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly
225                 230                 235                 240

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
                245                 250                 255
```

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser
260 265 270

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr
275 280 285

Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
290 295 300

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser
305 310 315 320

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr
325 330 335

Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr
340 345 350

Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser
355 360 365

Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser
370 375 380

Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
385 390 395

<210> 162
<211> 432
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 162
Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
1 5 10 15

Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr
20 25 30

Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser
35 40 45

Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
50 55 60

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser
65 70 75 80

```
Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr
                85                  90                  95

Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
            100                 105                 110

Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala
        115                 120                 125

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
    130                 135                 140

Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
145                 150                 155                 160

Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr
                165                 170                 175

Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser
            180                 185                 190

Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
        195                 200                 205

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr
    210                 215                 220

Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser
225                 230                 235                 240

Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
                245                 250                 255

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser
            260                 265                 270

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr
        275                 280                 285

Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
    290                 295                 300

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser
305                 310                 315                 320

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala
                325                 330                 335

Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
            340                 345                 350
```

Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser
355 360 365

Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala
370 375 380

Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
385 390 395 400

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr
405 410 415

Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser
420 425 430

<210> 163
<211> 468
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 163
Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
1 5 10 15

Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
20 25 30

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
35 40 45

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
50 55 60

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
65 70 75 80

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly
85 90 95

Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
100 105 110

Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
115 120 125

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
130 135 140

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
145 150 155 160

```
Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
                165                 170                 175

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser
            180                 185                 190

Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
        195                 200                 205

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
    210                 215                 220

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
225                 230                 235                 240

Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
                245                 250                 255

Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
            260                 265                 270

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
        275                 280                 285

Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
    290                 295                 300

Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
305                 310                 315                 320

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser
                325                 330                 335

Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
            340                 345                 350

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
        355                 360                 365

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
    370                 375                 380

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
385                 390                 395                 400

Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
                405                 410                 415

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
            420                 425                 430
```

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
435 440 445

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
450 455 460

Ala Ser Pro Gly
465

<210> 164
<211> 504
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 164
Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
1 5 10 15

Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
20 25 30

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
35 40 45

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
50 55 60

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
65 70 75 80

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly
85 90 95

Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
100 105 110

Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
115 120 125

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
130 135 140

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
145 150 155 160

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
165 170 175

```
Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser
            180                 185                 190

Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
        195                 200                 205

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
    210                 215                 220

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
225                 230                 235                 240

Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
                245                 250                 255

Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
            260                 265                 270

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
        275                 280                 285

Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
    290                 295                 300

Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
305                 310                 315                 320

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser
                325                 330                 335

Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
            340                 345                 350

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
        355                 360                 365

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
    370                 375                 380

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
385                 390                 395                 400

Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
                405                 410                 415

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
            420                 425                 430

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
        435                 440                 445
```

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
450 455 460

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser
465 470 475 480

Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
485 490 495

Ser Ser Pro Gly Ser Ser Thr Pro
500

<210> 165
<211> 540
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 165
Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
1 5 10 15

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
20 25 30

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly
35 40 45

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
50 55 60

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65 70 75 80

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
85 90 95

Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
100 105 110

Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
115 120 125

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
130 135 140

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145 150 155 160

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
165 170 175

```
Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
            180                 185                 190

Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
        195                 200                 205

Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
    210                 215                 220

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
225                 230                 235                 240

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
                245                 250                 255

Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
            260                 265                 270

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser
        275                 280                 285

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
    290                 295                 300

Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
305                 310                 315                 320

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
                325                 330                 335

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
            340                 345                 350

Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
        355                 360                 365

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser
    370                 375                 380

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
385                 390                 395                 400

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
                405                 410                 415

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly
            420                 425                 430

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
        435                 440                 445
```

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
450 455 460

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly
465 470 475 480

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
485 490 495

Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
500 505 510

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
515 520 525

Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly
530 535 540

<210> 166
<211> 576
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 166
Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
1 5 10 15

Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
20 25 30

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
35 40 45

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
50 55 60

Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
65 70 75 80

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
85 90 95

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
100 105 110

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
115 120 125

```
Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly
    130                 135                 140

Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
145                 150                 155                 160

Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
                165                 170                 175

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
            180                 185                 190

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
        195                 200                 205

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
    210                 215                 220

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser
225                 230                 235                 240

Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
                245                 250                 255

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
            260                 265                 270

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
        275                 280                 285

Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
    290                 295                 300

Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
305                 310                 315                 320

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
                325                 330                 335

Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
            340                 345                 350

Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
        355                 360                 365

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser
    370                 375                 380

Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
385                 390                 395                 400
```

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
405 410 415

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
420 425 430

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
435 440 445

Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
450 455 460

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
465 470 475 480

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
485 490 495

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
500 505 510

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser
515 520 525

Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
530 535 540

Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
545 550 555 560

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly
565 570 575

<210> 167
<211> 612
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 167
Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
1 5 10 15

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser
20 25 30

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
35 40 45

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
50 55 60

```
Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser
65                  70                  75                  80

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly
                85                  90                  95

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
            100                 105                 110

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
        115                 120                 125

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
    130                 135                 140

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
145                 150                 155                 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
                165                 170                 175

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
            180                 185                 190

Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
        195                 200                 205

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser
    210                 215                 220

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
225                 230                 235                 240

Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
                245                 250                 255

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
            260                 265                 270

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
        275                 280                 285

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
    290                 295                 300

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
305                 310                 315                 320

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly
                325                 330                 335
```

```
Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
            340                 345                 350

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
        355                 360                 365

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
    370                 375                 380

Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
385                 390                 395                 400

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
                405                 410                 415

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
            420                 425                 430

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
        435                 440                 445

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro
    450                 455                 460

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
465                 470                 475                 480

Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
                485                 490                 495

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
            500                 505                 510

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
        515                 520                 525

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
    530                 535                 540

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
545                 550                 555                 560

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
                565                 570                 575

Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
            580                 585                 590

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
        595                 600                 605
```

Pro Gly Thr Ser
610

<210> 168
<211> 648
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 168
Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
1 5 10 15

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
20 25 30

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro
35 40 45

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
50 55 60

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65 70 75 80

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
85 90 95

Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
100 105 110

Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
115 120 125

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
130 135 140

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
145 150 155 160

Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
165 170 175

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
180 185 190

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
195 200 205

```
Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
    210                 215                 220

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly
225                 230                 235                 240

Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
                245                 250                 255

Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
            260                 265                 270

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
        275                 280                 285

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
    290                 295                 300

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
305                 310                 315                 320

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser
                325                 330                 335

Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
            340                 345                 350

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
        355                 360                 365

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
    370                 375                 380

Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
385                 390                 395                 400

Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
                405                 410                 415

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
            420                 425                 430

Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
        435                 440                 445

Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
    450                 455                 460

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser
465                 470                 475                 480
```

Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
485 490 495

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
500 505 510

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
515 520 525

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
530 535 540

Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
545 550 555 560

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
565 570 575

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
580 585 590

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
595 600 605

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser
610 615 620

Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
625 630 635 640

Ser Ser Pro Gly Ser Ser Thr Pro
645

<210> 169
<211> 684
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 169
Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
1 5 10 15

Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
20 25 30

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser
35 40 45

Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
50 55 60

```
Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
65                  70                  75                  80

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
                85                  90                  95

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
            100                 105                 110

Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
        115                 120                 125

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
    130                 135                 140

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
145                 150                 155                 160

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
                165                 170                 175

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly
            180                 185                 190

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
        195                 200                 205

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
    210                 215                 220

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
225                 230                 235                 240

Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
                245                 250                 255

Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
            260                 265                 270

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
        275                 280                 285

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
    290                 295                 300

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
305                 310                 315                 320

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
                325                 330                 335
```

```
Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
            340                 345                 350

Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
        355                 360                 365

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
    370                 375                 380

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
385                 390                 395                 400

Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
                405                 410                 415

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser
            420                 425                 430

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
        435                 440                 445

Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
    450                 455                 460

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
465                 470                 475                 480

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
                485                 490                 495

Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
            500                 505                 510

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser
        515                 520                 525

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
    530                 535                 540

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
545                 550                 555                 560

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly
                565                 570                 575

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
            580                 585                 590

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
        595                 600                 605
```

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly
610 615 620

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
625 630 635 640

Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
645 650 655

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
660 665 670

Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly
675 680

<210> 170
<211> 720
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 170
Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
1 5 10 15

Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
20 25 30

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser
35 40 45

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
50 55 60

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
65 70 75 80

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro
85 90 95

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
100 105 110

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
115 120 125

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
130 135 140

```
Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
145                 150                 155                 160

Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
                165                 170                 175

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
            180                 185                 190

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
        195                 200                 205

Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
    210                 215                 220

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
225                 230                 235                 240

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
                245                 250                 255

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
            260                 265                 270

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly
        275                 280                 285

Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
    290                 295                 300

Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
305                 310                 315                 320

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
                325                 330                 335

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
            340                 345                 350

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
        355                 360                 365

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser
    370                 375                 380

Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
385                 390                 395                 400

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
                405                 410                 415
```

```
Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
            420                 425                 430

Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
        435                 440                 445

Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
    450                 455                 460

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
465                 470                 475                 480

Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
                485                 490                 495

Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
            500                 505                 510

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser
        515                 520                 525

Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
    530                 535                 540

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
545                 550                 555                 560

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
                565                 570                 575

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
            580                 585                 590

Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
        595                 600                 605

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
    610                 615                 620

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
625                 630                 635                 640

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
                645                 650                 655

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser
            660                 665                 670

Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
        675                 680                 685
```

Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
690 695 700

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly
705 710 715 720

<210> 171
<211> 756
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 171
Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
1 5 10 15

Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
20 25 30

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
35 40 45

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
50 55 60

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65 70 75 80

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
85 90 95

Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
100 105 110

Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
115 120 125

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser
130 135 140

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145 150 155 160

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
165 170 175

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
180 185 190

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
195 200 205

```
Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
    210                 215                 220

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
225                 230                 235                 240

Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
                245                 250                 255

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
            260                 265                 270

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser
        275                 280                 285

Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
    290                 295                 300

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
305                 310                 315                 320

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser
                325                 330                 335

Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
            340                 345                 350

Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
        355                 360                 365

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
    370                 375                 380

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
385                 390                 395                 400

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
                405                 410                 415

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser
            420                 425                 430

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
        435                 440                 445

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
    450                 455                 460

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser
465                 470                 475                 480
```

```
Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
                485                 490                 495

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
            500                 505                 510

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly
        515                 520                 525

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
    530                 535                 540

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
545                 550                 555                 560

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser
                565                 570                 575

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
            580                 585                 590

Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
        595                 600                 605

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser
    610                 615                 620

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
625                 630                 635                 640

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
                645                 650                 655

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser
            660                 665                 670

Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
        675                 680                 685

Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
    690                 695                 700

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser
705                 710                 715                 720

Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
                725                 730                 735

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
            740                 745                 750
```

Ala Ser Pro Gly
755

<210> 172
<211> 792
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 172
Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
1 5 10 15

Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
20 25 30

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
35 40 45

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
50 55 60

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65 70 75 80

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
85 90 95

Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
100 105 110

Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
115 120 125

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser
130 135 140

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145 150 155 160

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
165 170 175

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
180 185 190

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
195 200 205

```
Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
    210                 215                 220

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
225                 230                 235                 240

Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
                245                 250                 255

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
            260                 265                 270

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser
        275                 280                 285

Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
    290                 295                 300

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
305                 310                 315                 320

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser
                325                 330                 335

Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
            340                 345                 350

Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
        355                 360                 365

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
    370                 375                 380

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
385                 390                 395                 400

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
                405                 410                 415

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser
            420                 425                 430

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
        435                 440                 445

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
    450                 455                 460

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser
465                 470                 475                 480
```

```
Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
                485                 490                 495

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
            500                 505                 510

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly
        515                 520                 525

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
    530                 535                 540

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
545                 550                 555                 560

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser
                565                 570                 575

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
            580                 585                 590

Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
        595                 600                 605

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser
    610                 615                 620

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
625                 630                 635                 640

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
                645                 650                 655

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser
            660                 665                 670

Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
        675                 680                 685

Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
    690                 695                 700

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser
705                 710                 715                 720

Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
                725                 730                 735

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
            740                 745                 750
```

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro
755 760 765

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
770 775 780

Thr Gly Pro Gly Ala Ser Pro Gly
785 790

<210> 173
<211> 828
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 173
Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
1 5 10 15

Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
20 25 30

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
35 40 45

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
50 55 60

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65 70 75 80

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
85 90 95

Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
100 105 110

Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
115 120 125

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser
130 135 140

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145 150 155 160

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
165 170 175

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
180 185 190

```
Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
        195                 200                 205

Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
    210                 215                 220

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
225                 230                 235                 240

Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
                245                 250                 255

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
            260                 265                 270

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser
        275                 280                 285

Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
    290                 295                 300

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
305                 310                 315                 320

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser
                325                 330                 335

Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
            340                 345                 350

Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
        355                 360                 365

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
    370                 375                 380

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
385                 390                 395                 400

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
                405                 410                 415

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser
            420                 425                 430

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
        435                 440                 445

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
    450                 455                 460
```

```
Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser
465                 470                 475                 480

Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
                485                 490                 495

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
            500                 505                 510

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly
        515                 520                 525

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
    530                 535                 540

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
545                 550                 555                 560

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser
                565                 570                 575

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
            580                 585                 590

Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
        595                 600                 605

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser
    610                 615                 620

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
625                 630                 635                 640

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
                645                 650                 655

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser
            660                 665                 670

Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
        675                 680                 685

Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
    690                 695                 700

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser
705                 710                 715                 720

Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
                725                 730                 735
```

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
740 745 750

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro
755 760 765

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
770 775 780

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
785 790 795 800

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser
805 810 815

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
820 825

<210> 174
<211> 4
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 174
Gly Ser Pro Gly
1

<210> 175
<211> 4
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 175
Glu Thr Glu Thr
1

<210> 176
<211> 5
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 176
Pro Gly Ser Ser Ser
1 5

<210> 177
<211> 5
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 177
Gly Pro Ser Gly Pro
1 5

<210> 178
<211> 10
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 178
Gly Ala Gly Ser Pro Gly Ala Glu Thr Ala
1 5 10

<210> 179
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 179
Lys Leu Thr Arg Ala Glu Thr
1 5

<210> 180
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 180
Asp Phe Thr Arg Val Val Gly
1 5

<210> 181
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 181
Thr Met Thr Arg Ile Val Gly Gly
1 5

<210> 182
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 182
Ser Pro Phe Arg Ser Thr Gly Gly
1 5

<210> 183
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 183
Leu Gln Val Arg Ile Val Gly Gly
1 5

<210> 184
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 184
Pro Leu Gly Arg Ile Val Gly Gly
1 5

<210> 185
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 185
Ile Glu Gly Arg Thr Val Gly Gly
1 5

<210> 186
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 186

Leu Thr Pro Arg Ser Leu Leu Val
1 5

<210> 187
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 187
Leu Gly Pro Val Ser Gly Val Pro
1 5

<210> 188
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 188
Val Ala Gly Asp Ser Leu Glu Glu
1 5

<210> 189
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 189
Gly Pro Ala Gly Leu Gly Gly Ala
1 5

<210> 190
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<220>
<221> MOD_RES
<222> (2)..(2)
<223> Pro or Ala

<220>
<221> MOD_RES
<222> (3)..(3)
<223> Any amino acid

<220>
<221> MOD_RES

<222> (6)..(6)
<223> Any amino acid

<220>
<221> MOD_RES
<222> (8)..(8)
<223> Any amino acid

<400> 190
Gly Xaa Xaa Gly Leu Xaa Gly Xaa
1   5

<210> 191
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
      peptide

<400> 191
Gly Pro Ala Gly Leu Arg Gly Ala
1   5

<210> 192
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
      peptide

<220>
<221> MOD_RES
<222> (3)..(3)
<223> Any amino acid

<220>
<221> MOD_RES
<222> (6)..(6)
<223> Any amino acid

<220>
<221> MOD_RES
<222> (7)..(7)
<223> Gly or Ala

<220>
<221> MOD_RES
<222> (8)..(8)
<223> Any amino acid

<400> 192
Gly Pro Xaa Gly Leu Xaa Xaa Xaa
1   5

<210> 193
<211> 8
<212> PRT
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic peptide

<400> 193
Ala Pro Leu Gly Leu Arg Leu Arg
1 5

<210> 194
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 194
Pro Ala Leu Pro Leu Val Ala Gln
1 5

<210> 195
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 195
Glu Asn Leu Tyr Phe Gln Gly
1 5

<210> 196
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 196
Glu Asn Leu Tyr Phe Gln Gly Ser
1 5

<210> 197
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 197
Asp Asp Asp Lys Ile Val Gly Gly
1 5

<210> 198
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 198
Asp Asp Asp Lys Ile Val Gly Gly
1 5

<210> 199
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 199
Leu Glu Val Leu Phe Gln Gly Pro
1 5

<210> 200
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 200
Leu Glu Val Leu Phe Gln Gly Pro
1 5

<210> 201
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 201
Leu Pro Lys Thr Gly Ser Glu Ser
1 5

<210> 202
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<220>
<221> MOD_RES
<222> (3)..(3)
<223> Lys, Glu, Ala or Asp

<220>

<221> MOD_RES
<222> (6)..(6)
<223> Any amino acid

<220>
<221> MOD_RES
<222> (7)..(7)
<223> Glu, Lys or Ser

<400> 202
Leu Pro Xaa Thr Gly Xaa Xaa Ser
1 5

<210> 203
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 203
Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Glu Ser Pro
1 5 10 15

Gly Gly Ser Ser Gly Ser Glu Ser Gly Glu Ser Pro Gly Gly Ser Ser
20 25 30

Gly Ser Glu Ser
35

<210> 204
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 204
ggttctggtg gcgaaccgtc cgagtctggt agctcaggtg aatctccggg tggctctagc 60

ggttccgagt caggtgaatc tcctggtggt tccagcggtt ccgagtca 108

<210> 205
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 205
Gly Ser Glu Gly Ser Ser Gly Pro Gly Glu Ser Ser Gly Glu Ser Pro
1 5 10 15

Gly Gly Ser Ser Gly Ser Glu Ser Gly Ser Ser Glu Ser Gly Ser Ser
20 25 30

Glu Gly Gly Pro
35

<210> 206
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 206
ggtagcgaag gttcttctgg tcctggcgag tcttcaggtg aatctcctgg tggttccagc 60

ggttctgaat caggttcctc cgaaagcggt tcttccgagg gcggtcca 108

<210> 207
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 207
Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Ser Glu
1 5 10 15

Ser Gly Ser Ser Glu Gly Gly Pro Gly Glu Ser Pro Gly Gly Ser Ser
20 25 30

Gly Ser Glu Ser
35

<210> 208
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 208
ggttcctctg aaagcggttc ttccgaaggt ggtccaggtt cctctgaaag cggttcttct 60

gagggtggtc caggtgaatc tccgggtggc tccagcggtt ccgagtca 108

<210> 209
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 209
Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Ser Glu

1 5 10 15

Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Gly Gly Glu Pro Ser Glu
20 25 30

Ser Gly Ser Ser
35

<210> 210
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 210
ggttccggtg gcgaaccgtc tgaatctggt agctcaggtt cttctgaaag cggttcttcc 60

gagggtggtc caggttctgg tggtgaacct tccgagtctg gtagctca 108

<210> 211
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 211
Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Glu Gly
1 5 10 15

Ser Ser Gly Pro Gly Glu Ser Ser Gly Ser Glu Gly Ser Ser Gly Pro
20 25 30

Gly Glu Ser Ser
35

<210> 212
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 212
ggttcttccg aaagcggttc ttctgagggt ggtccaggta gcgaaggttc ttccggtcca 60

ggtgagtctt caggtagcga aggttcttct ggtcctggtg aatcttca 108

<210> 213
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 213
Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro Gly Glu Ser Pro
1 5 10 15

Gly Gly Ser Ser Gly Ser Glu Ser Gly Ser Glu Gly Ser Ser Gly Pro
20 25 30

Gly Glu Ser Ser
35

<210> 214
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 214
ggttcctctg aaagcggttc ttccgagggt ggtccaggtg aatctccagg tggttccagc 60

ggttctgagt caggtagcga aggttcttct ggtccaggtg aatcctca 108

<210> 215
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 215
Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Gly Gly
1 5 10 15

Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Glu Gly Ser Ser Gly Pro
20 25 30

Gly Glu Ser Ser
35

<210> 216
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 216
ggttctggtg gtgaaccgtc tgagtctggt agctcaggtt ccggtggcga accatccgaa 60

tctggtagct caggtagcga aggttcttcc ggtccaggtg agtcttca 108

<210> 217
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 217
Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Glu Gly
1 5 10 15

Ser Ser Gly Pro Gly Glu Ser Ser Gly Glu Ser Pro Gly Gly Ser Ser
20 25 30

Gly Ser Glu Ser
35

<210> 218
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 218
ggttcttccg aaagcggttc ttccgaaggc ggtccaggta gcgaaggttc ttctggtcca 60

ggcgaatctt caggtgaatc tcctggtggc tccagcggtt ctgagtca 108

<210> 219
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 219
Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Ser Glu
1 5 10 15

Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Ser Glu Ser Gly Ser Ser
20 25 30

Glu Gly Gly Pro
35

<210> 220
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 220
ggttcctccg aaagcggttc ttctgagggc ggtccaggtt cctccgaaag cggttcttcc 60

gagggcggtc caggttcttc tgaaagcggt tcttccgagg gcggtcca 108

<210> 221
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 221
Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Glu Gly
1 5 10 15

Ser Ser Gly Pro Gly Glu Ser Ser Gly Ser Ser Glu Ser Gly Ser Ser
20 25 30

Glu Gly Gly Pro
35

<210> 222
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 222
ggttccggtg gcgaaccgtc cgaatctggt agctcaggta gcgaaggttc ttctggtcca 60

ggcgaatctt caggttcctc tgaaagcggt tcttctgagg gcggtcca 108

<210> 223
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 223
Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Ser Glu
1 5 10 15

Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Gly Gly Glu Pro Ser Glu
20 25 30

Ser Gly Ser Ser
35

<210> 224
<211> 108

<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 224
ggttctggtg gtgaaccgtc cgaatctggt agctcaggtt cttccgaaag cggttcttct 60

gaaggtggtc caggttccgg tggcgaacct tctgaatctg gtagctca 108

<210> 225
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 225
Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Ser Glu
1 5 10 15

Ser Gly Ser Ser Glu Gly Gly Pro Gly Glu Ser Pro Gly Gly Ser Ser
20 25 30

Gly Ser Glu Ser
35

<210> 226
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 226
ggttctggtg gcgaaccgtc cgaatctggt agctcaggtt cctccgaaag cggttcttct 60

gaaggtggtc caggtgaatc tccaggtggt tctagcggtt ctgaatca 108

<210> 227
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 227
Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Glu Ser Pro
1 5 10 15

Gly Gly Ser Ser Gly Ser Glu Ser Gly Ser Glu Gly Ser Ser Gly Pro
20 25 30

Gly Glu Ser Ser
35

<210> 228
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 228
ggttctggtg gcgaaccgtc tgagtctggt agctcaggtg aatctcctgg tggctccagc 60

ggttctgaat caggtagcga aggttcttct ggtcctggtg aatcttca 108

<210> 229
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 229
Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Glu Ser Pro
1 5 10 15

Gly Gly Ser Ser Gly Ser Glu Ser Gly Ser Gly Gly Glu Pro Ser Glu
20 25 30

Ser Gly Ser Ser
35

<210> 230
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 230
ggttccggtg gcgaaccttc cgaatctggt agctcaggtg aatctccggg tggttctagc 60

ggttctgagt caggttctgg tggtgaacct tccgagtctg gtagctca 108

<210> 231
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 231
Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Ser Glu
1 5 10 15

Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Ser Glu Ser Gly Ser Ser
20 25 30

Glu Gly Gly Pro
35

<210> 232
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 232
ggttcctctg aaagcggttc ttctgagggc ggtccaggtt cttccgaaag cggttcttcc 60

gagggcggtc caggttcttc cgaaagcggt tcttctgaag gcggtcca 108

<210> 233
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 233
Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser Gly Ser Glu Gly
1 5 10 15

Ser Ser Gly Pro Gly Glu Ser Ser Gly Ser Glu Gly Ser Ser Gly Pro
20 25 30

Gly Glu Ser Ser
35

<210> 234
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 234
ggtgaatctc cgggtggctc cagcggttct gagtcaggta gcgaaggttc ttccggtccg 60

ggtgagtcct caggtagcga aggttcttcc ggtcctggtg agtcttca 108

<210> 235
<211> 36
<212> PRT
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 235
Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Gly Gly
1 5 10 15

Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Ser Glu Ser Gly Ser Ser
20 25 30

Glu Gly Gly Pro
35

<210> 236
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 236
ggttctggtg gcgaaccttc cgaatctggt agctcaggtt ccggtggtga accttctgaa 60

tctggtagct caggttcttc tgaaagcggt tcttccgagg gcggtcca 108

<210> 237
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 237
Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Gly Gly
1 5 10 15

Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Gly Gly Glu Pro Ser Glu
20 25 30

Ser Gly Ser Ser
35

<210> 238
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 238
ggttccggtg gtgaaccttc tgaatctggt agctcaggtt ccggtggcga accatccgag 60

tctggtagct caggttccgg tggtgaacca tccgagtctg gtagctca 108

<210> 239
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 239
Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Ser Glu
1 5 10 15

Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Glu Gly Ser Ser Gly Pro
20 25 30

Gly Glu Ser Ser
35

<210> 240
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 240
ggttccggtg gcgaaccttc tgaatctggt agctcaggtt cctccgaaag cggttcttct 60

gagggcggtc caggtagcga aggttcttct ggtccgggcg agtcttca 108

<210> 241
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 241
Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Glu Gly
1 5 10 15

Ser Ser Gly Pro Gly Glu Ser Ser Gly Ser Gly Gly Glu Pro Ser Glu
20 25 30

Ser Gly Ser Ser
35

<210> 242
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 242
ggttccggtg gtgaaccgtc cgagtctggt agctcaggta gcgaaggttc ttctggtccg 60

ggtgagtctt caggttctgg tggcgaaccg tccgaatctg gtagctca 108

<210> 243
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 243
Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Glu Ser Pro
1 5 10 15

Gly Gly Ser Ser Gly Ser Glu Ser Gly Ser Gly Gly Glu Pro Ser Glu
20 25 30

Ser Gly Ser Ser
35

<210> 244
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 244
ggttctggtg gcgaaccgtc cgaatctggt agctcaggtg aatctcctgg tggttccagc 60

ggttccgagt caggttctgg tggcgaacct tccgaatctg gtagctca 108

<210> 245
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 245
Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Gly Gly
1 5 10 15

Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Ser Glu Ser Gly Ser Ser
20 25 30

Glu Gly Gly Pro
35

<210> 246
<211> 108
<212> DNA

<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 246
ggttcttccg aaagcggttc ttccgagggc ggtccaggtt ccggtggtga accatctgaa 60

tctggtagct caggttcttc tgaaagcggt tcttctgaag gtggtcca 108

<210> 247
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 247
Gly Ser Glu Gly Ser Ser Gly Pro Gly Glu Ser Ser Gly Glu Ser Pro
1 5 10 15

Gly Gly Ser Ser Gly Ser Glu Ser Gly Ser Glu Gly Ser Ser Gly Pro
20 25 30

Gly Glu Ser Ser
35

<210> 248
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 248
ggtagcgaag gttcttccgg tcctggtgag tcttcaggtg aatctccagg tggctctagc 60

ggttccgagt caggtagcga aggttcttct ggtcctggcg agtcctca 108

<210> 249
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 249
Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Ser Glu
1 5 10 15

Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Ser Glu Ser Gly Ser Ser
20 25 30

Glu Gly Gly Pro

35

<210> 250
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 250
ggttcctctg aaagcggttc ttccgaaggc ggtccaggtt cttccgaaag cggttcttct 60

gagggcggtc caggttcctc cgaaagcggt tcttctgagg gtggtcca 108

<210> 251
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 251
Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Glu Ser Pro
1 5 10 15

Gly Gly Ser Ser Gly Ser Glu Ser Gly Glu Ser Pro Gly Gly Ser Ser
20 25 30

Gly Ser Glu Ser
35

<210> 252
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 252
ggttctggtg gcgaaccttc cgagtctggt agctcaggtg aatctccggg tggttctagc 60

ggttccgagt caggtgaatc tccgggtggt tccagcggtt ctgagtca 108

<210> 253
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 253
Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Glu Gly
1 5 10 15

Ser Ser Gly Pro Gly Glu Ser Ser Gly Glu Ser Pro Gly Gly Ser Ser
20 25 30

Gly Ser Glu Ser
35

<210> 254
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 254
ggttctggtg gcgaaccatc tgagtctggt agctcaggta gcgaaggttc ttccggtcca 60

ggcgagtctt caggtgaatc tcctggtggc tccagcggtt ctgagtca 108

<210> 255
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 255
Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser Gly Glu Ser Pro
1 5 10 15

Gly Gly Ser Ser Gly Ser Glu Ser Gly Glu Ser Pro Gly Gly Ser Ser
20 25 30

Gly Ser Glu Ser
35

<210> 256
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 256
ggtgaatctc ctggtggttc agcggttcc gagtcaggtg aatctccagg tggctctagc 60

ggttccgagt caggtgaatc tcctggtggt tctagcggtt ctgaatca 108

<210> 257
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic

polypeptide

<400> 257
Gly Ser Glu Gly Ser Ser Gly Pro Gly Glu Ser Ser Gly Ser Glu Gly
1 5 10 15

Ser Ser Gly Pro Gly Glu Ser Ser Gly Ser Gly Gly Glu Pro Ser Glu
20 25 30

Ser Gly Ser Ser
35

<210> 258
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 258
ggtagcgaag gttcttccgg tccaggtgaa tcttcaggta gcgaaggttc ttctggtcct 60

ggtgaatcct caggttccgg tggcgaacca tctgaatctg gtagctca 108

<210> 259
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 259
Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Glu Gly
1 5 10 15

Ser Ser Gly Pro Gly Glu Ser Ser Gly Glu Ser Pro Gly Gly Ser Ser
20 25 30

Gly Ser Glu Ser
35

<210> 260
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 260
ggttctggtg gcgaaccatc cgaatctggt agctcaggta gcgaaggttc ttctggtcct 60

ggcgaatctt caggtgaatc tccaggtggc tctagcggtt ccgaatca 108

<210> 261

<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 261
Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Ser Glu
1 5 10 15

Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Gly Gly Glu Pro Ser Glu
20 25 30

Ser Gly Ser Ser
35

<210> 262
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 262
ggttccggtg gtgaaccgtc cgaatctggt agctcaggtt cctctgaaag cggttcttcc 60

gagggtggtc caggttccgg tggtgaacct tctgagtctg gtagctca 108

<210> 263
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 263
Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Gly Gly
1 5 10 15

Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Glu Gly Ser Ser Gly Pro
20 25 30

Gly Glu Ser Ser
35

<210> 264
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 264

ggttcctctg aaagcggttc ttctgagggc ggtccaggtt ctggtggcga accatctgaa 60

tctggtagct caggtagcga aggttcttcc ggtccgggtg aatcttca 108

<210> 265
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 265
Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Glu Gly
1 5 10 15

Ser Ser Gly Pro Gly Glu Ser Ser Gly Ser Glu Gly Ser Ser Gly Pro
20 25 30

Gly Glu Ser Ser
35

<210> 266
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 266
ggttctggtg gtgaaccgtc cgaatctggt agctcaggta gcgaaggttc ttctggtcct 60

ggcgagtctt caggtagcga aggttcttct ggtcctggtg aatcttca 108

<210> 267
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 267
Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Ser Glu
1 5 10 15

Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Gly Gly Glu Pro Ser Glu
20 25 30

Ser Gly Ser Ser
35

<210> 268
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 268
ggttctggtg gcgaaccatc cgagtctggt agctcaggtt cttccgaaag cggttcttcc 60

gaaggcggtc caggttctgg tggtgaaccg tccgaatctg gtagctca 108

<210> 269
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 269
Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Glu Ser Pro
1 5 10 15

Gly Gly Ser Ser Gly Ser Glu Ser Gly Glu Ser Pro Gly Gly Ser Ser
20 25 30

Gly Ser Glu Ser
35

<210> 270
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 270
ggttccggtg gcgaaccttc cgaatctggt agctcaggtg aatctccggg tggttctagc 60

ggttccgaat caggtgaatc tccaggtggt tctagcggtt ccgaatca 108

<210> 271
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 271
Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Gly Gly
1 5 10 15

Glu Pro Ser Glu Ser Gly Ser Ser Gly Glu Ser Pro Gly Gly Ser Ser
20 25 30

Gly Ser Glu Ser
35

<210> 272
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 272
ggttccggtg gtgaaccatc tgagtctggt agctcaggtt ccggtggcga accgtccgag 60

tctggtagct caggtgaatc tccgggtggt tccagcggtt ccgaatca 108

<210> 273
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 273
Gly Ser Glu Gly Ser Ser Gly Pro Gly Glu Ser Ser Gly Ser Ser Glu
1 5 10 15

Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Glu Gly Ser Ser Gly Pro
20 25 30

Gly Glu Ser Ser
35

<210> 274
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 274
ggtagcgaag gttcttctgg tccgggcgaa tcctcaggtt cctccgaaag cggttcttcc 60

gaaggtggtc caggtagcga aggttcttcc ggtcctggtg aatcttca 108

<210> 275
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 275
Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro Gly Glu Ser Pro
1 5 10 15

Gly Gly Ser Ser Gly Ser Glu Ser Gly Glu Ser Pro Gly Gly Ser Ser
20 25 30

Gly Ser Glu Ser
35

<210> 276
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 276
ggttcctctg aaagcggttc ttctgaaggc ggtccaggtg aatctccggg tggctccagc 60

ggttctgaat caggtgaatc tcctggtggc tccagcggtt ccgagtca 108

<210> 277
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 277
Gly Ser Glu Gly Ser Ser Gly Pro Gly Glu Ser Ser Gly Ser Glu Gly
1 5 10 15

Ser Ser Gly Pro Gly Glu Ser Ser Gly Ser Gly Gly Glu Pro Ser Glu
20 25 30

Ser Gly Ser Ser
35

<210> 278
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 278
ggtagcgaag gttcttctgg tccaggcgag tcttcaggta gcgaaggttc ttccggtcct 60

ggcgagtctt caggttccgg tggcgaaccg tccgaatctg gtagctca 108

<210> 279
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 279
Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro
35

<210> 280
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 280
ggtagcccgg caggctctcc gacctctact gaggaaggta cttctgaaag cgcaaccccg 60

gagtccggcc caggtacctc taccgaaccg tctgagggca gcgcacca 108

<210> 281
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 281
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
1 5 10 15

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro
35

<210> 282
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 282
ggtacttcta ccgaaccgtc cgaaggcagc gctccaggta cctctactga accttccgag 60

ggcagcgctc caggtacctc taccgaacct tctgaaggta gcgcacca 108

<210> 283
<211> 36

<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 283
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
20 25 30

Glu Ser Gly Pro
35

<210> 284
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 284
ggtacctcta ccgaaccgtc tgaaggtagc gcaccaggta cctctgaaag cgcaactcct 60

gagtccggtc caggtacttc tgaaagcgca accccggagt ctggccca 108

<210> 285
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 285
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
20 25 30

Glu Ser Gly Pro
35

<210> 286
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 286
ggtacttcta ctgaaccgtc tgaaggtagc gcaccaggta cttctgaaag cgcaaccccg 60

gaatccggcc caggtacctc tgaaagcgca accccggagt ccggccca 108

<210> 287
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 287
Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
20 25 30

Ser Thr Glu Glu
35

<210> 288
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 288
ggtagcgaac cggcaacctc cggctctgaa accccaggta cctctgaaag cgctactcct 60

gaatccggcc caggtagccc ggcaggttct ccgacttcca ctgaggaa 108

<210> 289
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 289
Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
1 5 10 15

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro
35

<210> 290
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 290
ggtacttctg aaagcgcaac ccctgaatcc ggtccaggta gcgaaccggc tacttctggc 60

tctgagactc caggtacttc taccgaaccg tccgaaggta gcgcacca 108

<210> 291
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 291
Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala
1 5 10 15

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Glu Pro Ala Thr Ser Gly
20 25 30

Ser Glu Thr Pro
35

<210> 292
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 292
ggtagcccgg ctggctctcc aacctccact gaggaaggta gcccggctgg ctctccaacc 60

tccactgaag aaggtagcga accggctacc tccggctctg aaactcca 108

<210> 293
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 293
Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro
35

<210> 294
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 294
ggtagcccgg ctggctctcc tacctctact gaggaaggta cttctgaaag cgctactcct 60

gagtctggtc caggtacctc tactgaaccg tccgaaggta gcgctcca 108

<210> 295
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 295
Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala
1 5 10 15

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro
35

<210> 296
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 296
ggtagccctg ctggctctcc gacttctact gaggaaggta gcccggctgg ttctccgact 60

tctactgagg aaggtacttc taccgaacct tccgaaggta gcgctcca 108

<210> 297
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 297
Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
1 5 10 15

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu

20 25 30

Gly Ser Ala Pro
35

<210> 298
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 298
ggtacttctg aaagcgctac tccggagtcc ggtccaggta cctctaccga accgtccgaa 60

ggcagcgctc caggtacttc tactgaacct tctgagggta gcgctcca 108

<210> 299
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 299
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
1 5 10 15

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro
35

<210> 300
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 300
ggtacctcta ccgaaccttc cgaaggtagc gctccaggta gcccggcagg ttctcctact 60

tccactgagg aaggtacttc taccgaacct tctgagggta gcgcacca 108

<210> 301
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 301
Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
1 5 10 15

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
20 25 30

Glu Ser Gly Pro
35

<210> 302
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 302
ggtagcgaac cggctacttc cggctctgag actccaggta gccctgctgg ctctccgacc 60

tctaccgaag aaggtacctc tgaaagcgct acccctgagt ctggccca 108

<210> 303
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 303
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
20 25 30

Glu Ser Gly Pro
35

<210> 304
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 304
ggtacttcta ccgaaccttc cgagggcagc gcaccaggta cttctgaaag cgctacccct 60

gagtccggcc caggtacttc tgaaagcgct actcctgaat ccggtcca 108

<210> 305
<211> 36
<212> PRT

<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 305
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
1 5 10 15

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
20 25 30

Ser Thr Glu Glu
35

<210> 306
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 306
ggtacttcta ctgaaccgtc tgaaggcagc gcaccaggta gcgaaccggc tacttccggt 60

tctgaaaccc caggtagccc agcaggttct ccaacttcta ctgaagaa 108

<210> 307
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 307
Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
20 25 30

Ser Glu Thr Pro
35

<210> 308
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 308
ggtagccctg ctggctctcc aacctccacc gaagaaggta cctctgaaag cgcaacccct 60

gaatccggcc caggtagcga accggcaacc tccggttctg aaacccca 108

<210> 309
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 309
Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
1 5 10 15

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
20 25 30

Ser Thr Glu Glu
35

<210> 310
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 310
ggtacttctg aaagcgctac tcctgagtcc ggcccaggta gcccggctgg ctctccgact 60

tccaccgagg aaggtagccc ggctggctct ccaacttcta ctgaagaa 108

<210> 311
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 311
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro
35

<210> 312
<211> 108
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 312
ggtacctcta ctgaaccttc tgagggcagc gctccaggta cttctgaaag cgctaccccg 60

gagtccggtc caggtacttc tactgaaccg tccgaaggta gcgcacca 108

<210> 313
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 313
Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
1 5 10 15

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
20 25 30

Ser Glu Thr Pro
35

<210> 314
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 314
ggtagcccgg caggctctcc gacttccacc gaggaaggta cctctactga accttctgag 60

ggtagcgctc caggtagcga accggcaacc tctggctctg aaacccca 108

<210> 315
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 315
Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
1 5 10 15

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro
35

<210> 316
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 316
ggtagcccag caggctctcc gacttccact gaggaaggta cttctactga accttccgaa 60

ggcagcgcac caggtacctc tactgaacct tctgagggca gcgctcca 108

<210> 317
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 317
Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
20 25 30

Ser Thr Glu Glu
35

<210> 318
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 318
ggtagcgaac ctgctacctc cggttctgaa accccaggta cctctgaaag cgcaactccg 60

gagtctggtc caggtagccc tgcaggttct cctacctcca ctgaggaa 108

<210> 319
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 319
Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
1 5 10 15

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro
35

<210> 320
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 320
ggtacctctg aaagcgctac tccggagtct ggcccaggta cctctactga accgtctgag 60

ggtagcgctc caggtacttc tactgaaccg tccgaaggta gcgcacca 108

<210> 321
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 321
Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
1 5 10 15

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro
35

<210> 322
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 322
ggtagcccgg ctggttctcc gacttccacc gaggaaggta cctctactga accttctgag 60

ggtagcgctc caggtacctc tactgaacct tccgaaggca gcgctcca 108

<210> 323
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 323

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
1 5 10 15

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro
35

<210> 324
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 324
ggtacttcta ccgaaccgtc cgagggcagc gctccaggta cttctactga accttctgaa 60

ggcagcgctc caggtacttc tactgaacct tccgaaggta gcgcacca 108

<210> 325
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 325
Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro
35

<210> 326
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 326
ggtagcgaac ctgcaacctc tggctctgaa accccaggta cctctgaaag cgctactcct 60

gaatctggcc caggtacttc tactgaaccg tccgagggca gcgcacca 108

<210> 327
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 327
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
1 5 10 15

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro
35

<210> 328
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 328
ggtacttcta ccgaaccgtc cgagggtagc gcaccaggta gcccagcagg ttctcctacc 60

tccaccgagg aaggtacttc taccgaaccg tccgagggta gcgcacca 108

<210> 329
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 329
Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
20 25 30

Ser Glu Thr Pro
35

<210> 330
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 330
ggtagcgaac cggcaacctc cggctctgaa actccaggta cttctgaaag cgctactccg 60

gaatccggcc caggtagcga accggctact tccggctctg aaacccca 108

<210> 331
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 331
Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
20 25 30

Ser Glu Thr Pro
35

<210> 332
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 332
ggtagcgaac cggcaacttc cggctctgaa accccaggta cttctgaaag cgctactcct 60

gagtctggcc caggtagcga acctgctacc tctggctctg aaacccca 108

<210> 333
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 333
Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
1 5 10 15

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro
35

<210> 334
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic

polynucleotide

<400> 334
ggtagcgaac cggcaacctc tggctctgaa actccaggta gcgaacctgc aacctccggc 60

tctgaaaccc caggtacttc tactgaacct tctgagggca gcgcacca 108

<210> 335
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 335
Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
1 5 10 15

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
20 25 30

Ser Glu Thr Pro
35

<210> 336
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 336
ggtacttctg aaagcgctac cccggaatct ggcccaggta gcgaaccggc tacttctggt 60

tctgaaaccc caggtagcga accggctacc tccggttctg aaactcca 108

<210> 337
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 337
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
1 5 10 15

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
20 25 30

Glu Ser Gly Pro
35

<210> 338

<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 338
ggtacctcta ctgaaccttc cgaaggcagc gctccaggta cctctaccga accgtccgag 60

ggcagcgcac caggtacttc tgaaagcgca acccctgaat ccggtcca 108

<210> 339
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 339
Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
20 25 30

Glu Ser Gly Pro
35

<210> 340
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 340
ggtagcgaac cggcaacctc tggctctgaa accccaggta cctctgaaag cgctactccg 60

gaatctggtc caggtacttc tgaaagcgct actccggaat ccggtcca 108

<210> 341
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 341
Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
1 5 10 15

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro
35

<210> 342
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 342
ggtagcgaac ctgctacctc cggctctgaa actccaggta gcgaaccggc tacttccggt 60

tctgaaactc caggtacctc taccgaacct tccgaaggca gcgcacca 108

<210> 343
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 343
Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr
1 5 10 15

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
20 25 30

Ser Glu Thr Pro
35

<210> 344
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 344
ggtagcgaac ctgctacttc tggttctgaa actccaggta cttctaccga accgtccgag 60

ggtagcgctc caggtagcga acctgctact tctggttctg aaactcca 108

<210> 345
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 345
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr

1 5 10 15

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
20 25 30

Ser Glu Thr Pro
35

<210> 346
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 346
ggtacctcta ccgaaccgtc cgagggtagc gcaccaggta cctctactga accgtctgag 60

ggtagcgctc caggtagcga accggcaacc tccggttctg aaactcca 108

<210> 347
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 347
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
1 5 10 15

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
20 25 30

Ser Thr Glu Glu
35

<210> 348
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 348
ggtacttcta ctgaaccttc cgaaggtagc gctccaggta gcgaacctgc tacttctggt 60

tctgaaaccc caggtagccc ggctggctct ccgacctcca ccgaggaa 108

<210> 349
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 349
Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
1 5 10 15

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
20 25 30

Glu Ser Gly Pro
35

<210> 350
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 350
ggtagcgaac cggctacttc cggctctgag actccaggta gcccagctgg ttctccaacc 60

tctactgagg aaggtacttc tgaaagcgct acccctgaat ctggtcca 108

<210> 351
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 351
Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
1 5 10 15

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
20 25 30

Glu Ser Gly Pro
35

<210> 352
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 352
ggtacctctg aaagcgcaac tcctgagtct ggcccaggta gcgaacctgc tacctccggc 60

tctgagactc caggtacctc tgaaagcgca accccggaat ctggtcca 108

<210> 353
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 353
Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Thr Ser Pro
1 5 10 15

Ser Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser
20 25 30

Ser Thr Ala Pro
35

<210> 354
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 354
ggtacttcta ctccggaaag cggttccgca tctccaggta cttctcctag cggtgaatct 60

tctactgctc caggtacctc tcctagcggc gaatcttcta ctgctcca 108

<210> 355
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 355
Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser
1 5 10 15

Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro Ser Gly Glu Ser
20 25 30

Ser Thr Ala Pro
35

<210> 356
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 356
ggtacttctc cgagcggtga atcttctacc gcaccaggtt ctactagctc taccgctgaa 60

tctccgggcc caggtacttc tccgagcggt gaatcttcta ctgctcca 108

<210> 357
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 357
Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro
1 5 10 15

Ser Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly
20 25 30

Ser Ala Ser Pro
35

<210> 358
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 358
ggttccacca gctctactgc tgaatctcct ggtccaggta cctctcctag cggtgaatct 60

tctactgctc caggtacttc tactcctgaa agcggctctg cttctcca 108

<210> 359
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 359
Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser
1 5 10 15

Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro Ser Gly Glu Ser
20 25 30

Ser Thr Ala Pro
35

<210> 360
<211> 108

<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 360
ggttctacca gctctactgc agaatctcct ggcccaggtt ccaccagctc taccgcagaa 60

tctccgggtc caggtacttc ccctagcggt gaatcttcta ccgcacca 108

<210> 361
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 361
Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro
1 5 10 15

Ser Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly
20 25 30

Ser Ala Ser Pro
35

<210> 362
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 362
ggttctacta gctctactgc tgaatctcct ggcccaggta cttctcctag cggtgaatct 60

tctaccgctc caggtacctc tactccggaa agcggttctg catctcca 108

<210> 363
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 363
Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Thr
1 5 10 15

Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser Glu Ser Pro Ser
20 25 30

Gly Thr Ala Pro
35

<210> 364
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 364
ggttctacca gctctaccgc agaatctcct ggtccaggta cctctactcc ggaaagcggc 60

tctgcatctc caggttctac tagcgaatct ccttctggca ctgcacca 108

<210> 365
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 365
Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Thr
1 5 10 15

Pro Glu Ser Gly Ser Ala Ser Pro Gly Thr Ser Thr Pro Glu Ser Gly
20 25 30

Ser Ala Ser Pro
35

<210> 366
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 366
ggttctacta gctctactgc agaatctcct ggcccaggta cctctactcc ggaaagcggc 60

tctgcatctc caggtacttc tacccctgaa agcggttctg catctcca 108

<210> 367
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 367
Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Pro
1 5 10 15

Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser
20 25 30

Gly Thr Ala Pro
35

<210> 368
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 368
ggttctacca gcgaatctcc ttctggcacc gctccaggta cctctcctag cggcgaatct 60

tctaccgctc caggttctac tagcgaatct ccttctggca ctgcacca 108

<210> 369
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 369
Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser
1 5 10 15

Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro Ser Gly Glu Ser
20 25 30

Ser Thr Ala Pro
35

<210> 370
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 370
ggttccacca gctctactgc agaatctccg ggcccaggtt ctactagctc tactgcagaa 60

tctccgggtc caggtacttc tcctagcggc gaatcttcta ccgctcca 108

<210> 371
<211> 36
<212> PRT
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 371
Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Thr
1 5 10 15

Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser Glu Ser Pro Ser
20 25 30

Gly Thr Ala Pro
35

<210> 372
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 372
ggttccacta gctctactgc agaatctcct ggcccaggta cctctacccc tgaaagcggc 60

tctgcatctc caggttctac cagcgaatcc ccgtctggca ccgcacca 108

<210> 373
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 373
Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser
1 5 10 15

Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro Ser Gly Glu Ser
20 25 30

Ser Thr Ala Pro
35

<210> 374
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 374
ggttctacta gctctactgc tgaatctccg ggtccaggtt ctaccagctc tactgctgaa 60

tctcctggtc caggtacctc cccgagcggt gaatcttcta ctgcacca 108

<210> 375
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 375
Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser
1 5 10 15

Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu
20 25 30

Ser Pro Gly Pro
35

<210> 376
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 376
ggttctacca gctctaccgc tgaatctcct ggcccaggtt ctaccagcga atccccgtct 60

ggcaccgcac caggttctac tagctctacc gctgaatctc cgggtcca 108

<210> 377
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 377
Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser
1 5 10 15

Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser Ser Thr Ala Glu
20 25 30

Ser Pro Gly Pro
35

<210> 378
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 378
ggttctacca gctctaccgc agaatctcct ggcccaggtt ccactagctc taccgctgaa 60

tctcctggtc caggttctac tagctctacc gctgaatctc ctggtcca 108

<210> 379
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 379
Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser
1 5 10 15

Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser Ser Thr Ala Glu
20 25 30

Ser Pro Gly Pro
35

<210> 380
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 380
ggttctacta gcgaatcccc ttctggtact gctccaggtt ccactagctc taccgctgaa 60

tctcctggcc caggttccac tagctctact gcagaatctc ctggtcca 108

<210> 381
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 381
Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser
1 5 10 15

Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser
20 25 30

Gly Thr Ala Pro
35

<210> 382
<211> 108
<212> DNA

<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 382
ggttctacta gcgaatctcc ttctggtacc gctccaggtt ctaccagcga atccccgtct 60

ggtactgctc caggttctac cagcgaatct ccttctggta ctgcacca 108

<210> 383
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 383
Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser
1 5 10 15

Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser Ser Thr Ala Glu
20 25 30

Ser Pro Gly Pro
35

<210> 384
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 384
ggttccacca gctctactgc tgaatctcct ggcccaggtt ctaccagctc tactgctgaa 60

tctccgggcc caggttccac cagctctacc gctgaatctc cgggtcca 108

<210> 385
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 385
Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser
1 5 10 15

Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro Ser Gly Glu Ser
20 25 30

Ser Thr Ala Pro

35

<210> 386
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 386
ggttccacta gctctaccgc agaatctcct ggtccaggtt ctactagctc tactgctgaa 60

tctccgggtc caggtacctc ccctagcggc gaatcttcta ccgctcca 108

<210> 387
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 387
Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr
1 5 10 15

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
20 25 30

Thr Gly Ser Pro
35

<210> 388
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 388
ggttctagcc cttctgcttc caccggtacc ggcccaggta gctctactcc gtctggtgca 60

actggctctc caggtagctc tactccgtct ggtgcaaccg gctcccca 108

<210> 389
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 389
Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Thr
1 5 10 15

Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser Ser Thr Ala Glu
20 25 30

Ser Pro Gly Pro
35

<210> 390
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 390
ggtacttccc ctagcggtga atcttctact gctccaggta cctctactcc ggaaagcggc 60

tccgcatctc caggttctac tagctctact gctgaatctc ctggtcca 108

<210> 391
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 391
Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser
1 5 10 15

Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Thr Pro Glu Ser Gly
20 25 30

Ser Ala Ser Pro
35

<210> 392
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 392
ggttctacta gctctaccgc tgaatctccg ggtccaggtt ctaccagctc tactgcagaa 60

tctcctggcc caggtacttc tactccggaa agcggttccg cttctcca 108

<210> 393
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic

polypeptide

<400> 393
Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser
1 5 10 15

Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Thr Pro Glu Ser Gly
20 25 30

Ser Ala Ser Pro
35

<210> 394
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 394
ggtacttctc ctagcggtga atcttctacc gctccaggtt ctaccagctc tactgctgaa 60

tctcctggcc caggtacttc taccccggaa agcggctccg cttctcca 108

<210> 395
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 395
Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser
1 5 10 15

Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser Ser Thr Ala Glu
20 25 30

Ser Pro Gly Pro
35

<210> 396
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 396
ggttctacta gcgaatcccc ttctggtact gcaccaggtt ctaccagctc tactgctgaa 60

tctccgggcc caggttccac cagctctacc gcagaatctc ctggtcca 108

<210> 397

<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 397
Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser
1 5 10 15

Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu
20 25 30

Ser Pro Gly Pro
35

<210> 398
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 398
ggttccacca gctctaccgc tgaatctccg ggcccaggtt ctaccagcga atccccttct 60

ggcactgcac caggttctac tagctctacc gcagaatctc cgggccca 108

<210> 399
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 399
Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro
1 5 10 15

Ser Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly
20 25 30

Ser Ala Ser Pro
35

<210> 400
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 400

ggttctacca gctctactgc tgaatctccg ggtccaggta cttccccgag cggtgaatct 60

tctactgcac caggtacttc tactccggaa agcggttccg cttctcca 108

<210> 401
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 401
Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser
1 5 10 15

Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser
20 25 30

Ser Thr Ala Pro
35

<210> 402
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 402
ggttctacca gcgaatctcc ttctggcacc gctccaggtt ctactagcga atccccgtct 60

ggtaccgcac caggtacttc tcctagcggc gaatcttcta ccgcacca 108

<210> 403
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 403
Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser
1 5 10 15

Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly
20 25 30

Ser Ala Ser Pro
35

<210> 404
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 404
ggttctacca gcgaatcccc ttctggtact gctccaggtt ctaccagcga atccccttct 60

ggcaccgcac caggtacttc tacccctgaa agcggctccg cttctcca 108

<210> 405
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 405
Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser
1 5 10 15

Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser Ser Thr Ala Glu
20 25 30

Ser Pro Gly Pro
35

<210> 406
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 406
ggtacctcta ctcctgaaag cggttctgca tctccaggtt ccactagctc taccgcagaa 60

tctccgggcc caggttctac tagctctact gctgaatctc ctggccca 108

<210> 407
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 407
Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser
1 5 10 15

Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser
20 25 30

Ser Thr Ala Pro
35

<210> 408
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 408
ggttctacca gcgaatcccc ttctggcact gcaccaggtt ctactagcga atccccttct 60

ggtaccgcac caggtacttc tccgagcggc gaatcttcta ctgctcca 108

<210> 409
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 409
Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser
1 5 10 15

Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser Glu Ser Pro Ser
20 25 30

Gly Thr Ala Pro
35

<210> 410
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 410
ggttctacta gctctaccgc tgaatctcct ggcccaggtt ccactagctc taccgcagaa 60

tctccgggcc caggttctac tagcgaatcc ccttctggta ccgctcca 108

<210> 411
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 411
Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser
1 5 10 15

Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Thr Pro Glu Ser Gly
20 25 30

Ser Ala Ser Pro
35

<210> 412
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 412
ggttccacca gctctactgc agaatctcct ggcccaggtt ctactagctc taccgcagaa 60

tctcctggtc caggtacctc tactcctgaa agcggttccg catctcca 108

<210> 413
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 413
Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser
1 5 10 15

Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser Glu Ser Pro Ser
20 25 30

Gly Thr Ala Pro
35

<210> 414
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 414
ggttctacta gctctactgc tgaatctccg ggcccaggtt ctactagctc taccgctgaa 60

tctccgggtc caggttctac tagcgaatct ccttctggta ccgctcca 108

<210> 415
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 415
Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser
1 5 10 15

Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu
20 25 30

Ser Pro Gly Pro
35

<210> 416
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 416
ggtacctccc cgagcggtga atcttctact gcaccaggtt ctactagcga atccccttct 60

ggtactgctc caggttccac cagctctact gcagaatctc cgggtcca 108

<210> 417
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 417
Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Pro
1 5 10 15

Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu
20 25 30

Ser Pro Gly Pro
35

<210> 418
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 418
ggttctacta gcgaatcccc gtctggtact gctccaggta cttcccctag cggtgaatct 60

tctactgctc caggttctac cagctctacc gcagaatctc cgggtcca 108

<210> 419
<211> 36

<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 419
Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser
1 5 10 15

Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser
20 25 30

Ser Thr Ala Pro
35

<210> 420
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 420
ggttctacca gctctaccgc tgaatctcct ggcccaggtt ctactagcga atctccgtct 60

ggcaccgcac caggtacttc ccctagcggt gaatcttcta ctgcacca 108

<210> 421
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 421
Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser
1 5 10 15

Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly
20 25 30

Ser Ala Ser Pro
35

<210> 422
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 422
ggttctacta gcgaatctcc ttctggcact gcaccaggtt ctaccagcga atctccgtct 60

ggcactgcac caggtacctc tacccctgaa agcggttccg cttctcca 108

<210> 423
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 423
Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser
1 5 10 15

Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu
20 25 30

Ser Pro Gly Pro
35

<210> 424
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 424
ggtacctcta ctccggaaag cggttccgca tctccaggtt ctaccagcga atccccgtct 60

ggcaccgcac caggttctac tagctctact gctgaatctc cgggccca 108

<210> 425
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 425
Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro
1 5 10 15

Ser Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser
20 25 30

Ser Thr Ala Pro
35

<210> 426
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 426
ggttctacta gctctactgc agaatctccg ggcccaggta cctctcctag cggtgaatct 60

tctaccgctc caggtacttc tccgagcggt gaatcttcta ccgctcca 108

<210> 427
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 427
Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Pro
1 5 10 15

Ser Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser
20 25 30

Ser Thr Ala Pro
35

<210> 428
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 428
ggtacctccc ctagcggcga atcttctact gctccaggta cctctcctag cggcgaatct 60

tctaccgctc caggtacctc ccctagcggt gaatcttcta ccgcacca 108

<210> 429
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 429
Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Thr
1 5 10 15

Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser Glu Ser Pro Ser
20 25 30

Gly Thr Ala Pro
35

<210> 430
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 430
ggttccacta gctctactgc tgaatctcct ggcccaggta cttctactcc ggaaagcggt 60

tccgcttctc caggttctac tagcgaatct ccgtctggca ccgcacca 108

<210> 431
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 431
Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Pro
1 5 10 15

Ser Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser
20 25 30

Ser Thr Ala Pro
35

<210> 432
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 432
ggttctacta gcgaatctcc gtctggcact gctccaggta cttctcctag cggtgaatct 60

tctaccgctc caggtacttc ccctagcggc gaatcttcta ccgctcca 108

<210> 433
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 433
Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Thr
1 5 10 15

Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser Ser Thr Ala Glu

20 25 30

Ser Pro Gly Pro
35

<210> 434
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 434
ggttctacta gcgaatctcc ttctggtacc gctccaggta cttctacccc tgaaagcggc 60

tccgcttctc caggttccac tagctctacc gctgaatctc cgggtcca 108

<210> 435
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 435
Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser
1 5 10 15

Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser Glu Ser Pro Ser
20 25 30

Gly Thr Ala Pro
35

<210> 436
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 436
ggttccacta gctctactgc tgaatctcct ggcccaggtt ctaccagctc taccgctgaa 60

tctcctggcc caggttctac cagcgaatct ccgtctggca ccgcacca 108

<210> 437
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 437
Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Thr
1 5 10 15

Pro Glu Ser Gly Ser Ala Ser Pro Gly Thr Ser Thr Pro Glu Ser Gly
20 25 30

Ser Ala Ser Pro
35

<210> 438
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 438
ggttctacta gcgaatcccc gtctggtacc gcaccaggta cttctacccc ggaaagcggc 60

tctgcttctc caggtacttc taccccggaa agcggctccg catctcca 108

<210> 439
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 439
Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Thr
1 5 10 15

Pro Glu Ser Gly Ser Ala Ser Pro Gly Thr Ser Thr Pro Glu Ser Gly
20 25 30

Ser Ala Ser Pro
35

<210> 440
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 440
ggttctacta gcgaatcccc gtctggtact gctccaggta cttctactcc tgaaagcggt 60

tccgcttctc caggtacctc tactccggaa agcggttctg catctcca 108

<210> 441
<211> 36
<212> PRT

<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 441
Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
1 5 10 15

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
20 25 30

Thr Gly Ser Pro
35

<210> 442
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 442
ggtgcatccc cgggcactag ctctaccggt tctccaggta ctcctggtag cggtactgct 60

tcttcttctc caggtagctc tactccttct ggtgctactg gttctcca 108

<210> 443
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 443
Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
1 5 10 15

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala
20 25 30

Thr Gly Ser Pro
35

<210> 444
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 444
ggtagctcta ccccttctgg tgctaccggc tctccaggtt ctagcccgtc tgcttctacc 60

ggtaccggtc caggtagctc tacccttct ggtgctactg gttctcca 108

<210> 445
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 445
Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
1 5 10 15

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala
20 25 30

Thr Gly Ser Pro
35

<210> 446
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 446
ggtgcatctc cgggtactag ctctaccggt tctccaggtt ctagcccttc tgcttccact 60

ggtaccggcc caggtagctc taccccgtct ggtgctactg gttcccca 108

<210> 447
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 447
Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
1 5 10 15

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
20 25 30

Thr Gly Ser Pro
35

<210> 448
<211> 108
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 448
ggtactccgg gcagcggtac tgcttcttcc tctccaggta gctctacccc ttctggtgca 60

actggttccc caggtgcatc ccctggtact agctctaccg gttctcca 108

<210> 449
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 449
Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
1 5 10 15

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Arg Pro Ser Ala Ser Thr
20 25 30

Gly Thr Gly Pro
35

<210> 450
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 450
ggtacccctg gcagcggtac tgcttcttct tctccaggtg cttcccctgg taccagctct 60

accggttctc caggttctag accttctgca tccaccggta ctggtcca 108

<210> 451
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 451
Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
1 5 10 15

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
20 25 30

Thr Gly Ser Pro
35

<210> 452
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 452
ggtgcatctc ctggtaccag ctctaccggt tctccaggta gctctactcc ttctggtgct 60

actggctctc caggtgcttc cccgggtacc agctctaccg gttctcca 108

<210> 453
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 453
Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
1 5 10 15

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
20 25 30

Thr Gly Ser Pro
35

<210> 454
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 454
ggtaccccgg gcagcggtac cgcatcttcc tctccaggta gctctacccc gtctggtgct 60

accggttccc caggtagctc taccccgtct ggtgcaaccg gctcccca 108

<210> 455
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 455
Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
1 5 10 15

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
20 25 30

Thr Gly Ser Pro
35

<210> 456
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 456
ggtgcatctc cgggcactag ctctactggt tctccaggtg catcccctgg cactagctct 60

actggttctc caggtgcttc tcctggtacc agctctactg gttctcca 108

<210> 457
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 457
Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
1 5 10 15

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
20 25 30

Thr Gly Ser Pro
35

<210> 458
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 458
ggtagctcta ctccgtctgg tgcaaccggc tccccaggtt ctagcccgtc tgcttccact 60

ggtactggcc caggtgcttc cccgggcacc agctctactg gttctcca 108

<210> 459
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 459

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
1 5 10 15

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
20 25 30

Ser Ser Ser Pro
35

<210> 460
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 460
ggtagctcta ctccttctgg tgctaccggt tccccaggta gctctactcc ttctggtgct 60

actggttccc caggtactcc gggcagcggt actgcttctt cctctcca 108

<210> 461
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 461
Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
1 5 10 15

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
20 25 30

Thr Gly Ser Pro
35

<210> 462
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 462
ggtagctcta ctccgtctgg tgcaaccggt tccccaggta gctctactcc ttctggtgct 60

actggctccc caggtgcatc ccctggcacc agctctaccg gttctcca 108

<210> 463
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 463
Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Pro
1 5 10 15

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala
20 25 30

Thr Gly Ser Pro
35

<210> 464
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 464
ggtactcctg gtagcggtac cgcatcttcc tctccaggtt ctagcccttc tgcatctacc 60

ggtaccggtc caggtagctc tactccttct ggtgctactg gctctcca 108

<210> 465
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 465
Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
1 5 10 15

Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala
20 25 30

Ser Ser Ser Pro
35

<210> 466
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 466
ggtgcttccc cgggcactag ctctaccggt tctccaggtt ctagcccttc tgcatctact 60

ggtactggcc caggtactcc gggcagcggt actgcttctt cctctcca 108

<210> 467
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 467
Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
1 5 10 15

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
20 25 30

Thr Gly Ser Pro
35

<210> 468
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 468
ggtagctcta ctccgtctgg tgctaccggc tctccaggta gctctacccc ttctggtgca 60

accggctccc caggtgcttc tccgggtacc agctctactg gttctcca 108

<210> 469
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 469
Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
1 5 10 15

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
20 25 30

Gly Thr Gly Pro
35

<210> 470
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic

polynucleotide

<400> 470
ggtacccctg gcagcggtac cgcttcttcc tctccaggta gctctacccc gtctggtgct 60

actggctctc caggttctag cccgtctgca tctaccggta ccggccca 108

<210> 471
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 471
Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
1 5 10 15

Ser Gly Thr Ala Ser Ser Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
20 25 30

Ser Ser Ser Pro
35

<210> 472
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 472
ggtagctcta ctccttctgg tgcaaccggc tccccaggta ccccgggcag cggtaccgca 60

tcttcctctc caggtactcc gggtagcggt actgcttctt cttctcca 108

<210> 473
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 473
Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
1 5 10 15

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
20 25 30

Thr Gly Ser Pro
35

<210> 474

<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 474
ggtaccccgg gtagcggtac tgcttcttcc tctccaggta gctctacccc ttctggtgca 60

accggctctc caggtgcttc tccgggcacc agctctaccg gttctcca 108

<210> 475
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 475
Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
1 5 10 15

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
20 25 30

Thr Gly Ser Pro
35

<210> 476
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 476
ggtagctcta ccccgtctgg tgctaccggc tctccaggta gctctacccc gtctggtgca 60

accggctccc caggtgcatc cccgggtact agctctaccg gttctcca 108

<210> 477
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 477
Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
1 5 10 15

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
20 25 30

Thr Gly Ser Pro
35

<210> 478
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 478
ggtgcttctc cgggcaccag ctctactggt tctccaggta ccccgggcag cggtaccgca 60

tcttcttctc caggtagctc tactccttct ggtgcaactg gttctcca 108

<210> 479
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 479
Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr
1 5 10 15

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
20 25 30

Ser Ser Ser Pro
35

<210> 480
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 480
ggttctagcc cgtctgcttc caccggtact ggcccaggta gctctacccc gtctggtgca 60

actggttccc caggtacccc tggtagcggt accgcttctt cttctcca 108

<210> 481
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 481
Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro

1 5 10 15

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala
20 25 30

Thr Gly Ser Pro
35

<210> 482
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 482
ggtgcttctc cgggcaccag ctctactggt tctccaggtt ctagcccttc tgcatccacc 60

ggtaccggtc caggtagctc tacccttct ggtgcaaccg gctctcca 108

<210> 483
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 483
Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
1 5 10 15

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
20 25 30

Thr Gly Ser Pro
35

<210> 484
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 484
ggtgcatccc cgggcaccag ctctaccggt tctccaggta gctctacccc gtctggtgct 60

accggctctc caggtagctc taccccgtct ggtgctactg gctctcca 108

<210> 485
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 485
Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
1 5 10 15

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
20 25 30

Ser Ser Ser Pro
35

<210> 486
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 486
ggtacccctg gtagcggtac tgcttcttcc tctccaggta gctctactcc gtctggtgct 60

accggttctc caggtacccc gggtagcggt accgcatctt cttctcca 108

<210> 487
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 487
Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr
1 5 10 15

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
20 25 30

Thr Gly Ser Pro
35

<210> 488
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 488
ggttctagcc cttctgcttc caccggtact ggcccaggta gctctacccc ttctggtgct 60

accggctccc caggtagctc tactccttct ggtgcaactg gctctcca 108

<210> 489
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 489
Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
1 5 10 15

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
20 25 30

Gly Thr Gly Pro
35

<210> 490
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 490
ggtgcttctc ctggcaccag ctctactggt tctccaggtt ctagcccttc tgcttctacc 60

ggtactggtc caggttctag cccttctgca tccactggta ctggtcca 108

<210> 491
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 491
Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
1 5 10 15

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
20 25 30

Thr Gly Ser Pro
35

<210> 492
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 492
ggtactcctg gcagcggtac cgcttcttct tctccaggtg cttctcctgg tactagctct 60

actggttctc caggtgcttc tccgggcact agctctactg gttctcca 108

<210> 493
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 493
Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro
1 5 10 15

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
20 25 30

Thr Gly Ser Pro
35

<210> 494
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 494
ggtagctcta ccccgtctgg tgctaccggt tccccaggtg cttctcctgg tactagctct 60

accggttctc caggtagctc taccccgtct ggtgctactg gctctcca 108

<210> 495
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 495
Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
1 5 10 15

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
20 25 30

Thr Gly Ser Pro
35

<210> 496
<211> 108

<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 496
ggtagctcta ccccgtctgg tgctactggt tctccaggta ctccgggcag cggtactgct 60

tcttcctctc caggtagctc taccccttct ggtgctactg gctctcca 108

<210> 497
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 497
Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
1 5 10 15

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala
20 25 30

Thr Gly Ser Pro
35

<210> 498
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 498
ggtgcatctc ctggtaccag ctctactggt tctccaggtt ctagcccttc tgcttctacc 60

ggtaccggtc caggtagctc tactccttct ggtgctaccg gttctcca 108

<210> 499
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 499
Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
1 5 10 15

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
20 25 30

Thr Gly Ser Pro
35

<210> 500
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 500
ggtagctcta ccccgtctgg tgctactggc tctccaggta gctctactcc ttctggtgct 60

actggttccc caggtagctc taccccgtct ggtgcaactg gctctcca 108

<210> 501
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 501
Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
1 5 10 15

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
20 25 30

Thr Gly Ser Pro
35

<210> 502
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 502
ggtgcatccc cgggtaccag ctctaccggt tctccaggta ctcctggcag cggtactgca 60

tcttcctctc caggtgcttc tccgggcacc agctctactg gttctcca 108

<210> 503
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 503
Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
1 5 10 15

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
20 25 30

Thr Gly Ser Pro
35

<210> 504
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 504
ggtagctcta ctccttctgg tgcaactggt tctccaggtt ctagcccgtc tgcatccact 60

ggtaccggtc caggtgcttc ccctggcacc agctctaccg gttctcca 108

<210> 505
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 505
Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
1 5 10 15

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
20 25 30

Gly Thr Gly Pro
35

<210> 506
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 506
ggtgcatctc ctggtactag ctctactggt tctccaggta gctctactcc gtctggtgca 60

accggctctc caggttctag cccttctgca tctaccggta ctggtcca 108

<210> 507
<211> 36
<212> PRT
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 507
Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
1 5 10 15

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
20 25 30

Thr Gly Ser Pro
35

<210> 508
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 508
ggtactcctg gcagcggtac cgcatcttcc tctccaggta gctctactcc gtctggtgca 60

actggttccc caggtgcttc tccgggtacc agctctaccg gttctcca 108

<210> 509
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 509
Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
1 5 10 15

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
20 25 30

Thr Gly Ser Pro
35

<210> 510
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 510
ggtagctcta ccccgtctgg tgcaaccggc tccccaggta ctcctggtag cggtaccgct 60

tcttcttctc caggtagctc tactccgtct ggtgctaccg gctcccca 108

<210> 511
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 511
Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
1 5 10 15

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
20 25 30

Thr Gly Ser Pro
35

<210> 512
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 512
ggttctagcc cttctgcatc caccggtacc ggcccaggtt ctagcccgtc tgcttctacc 60

ggtactggtc caggtgcttc tccgggtact agctctactg gttctcca 108

<210> 513
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 513
Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
1 5 10 15

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Asn Pro Ser Ala Ser Thr
20 25 30

Gly Thr Gly Pro
35

<210> 514
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 514
ggtaccccgg gtagcggtac cgcttcttct tctccaggta gctctactcc gtctggtgct 60

accggctctc caggttctaa cccttctgca tccaccggta ccggccca 108

<210> 515
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 515
Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr
1 5 10 15

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
20 25 30

Thr Gly Ser Pro
35

<210> 516
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 516
ggttctagcc cttctgcatc tactggtact ggcccaggta gctctactcc ttctggtgct 60

accggctctc caggtgcttc tccgggtact agctctaccg gttctcca 108

<210> 517
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 517
Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro
1 5 10 15

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
20 25 30

Ser Ser Ser Pro
35

<210> 518
<211> 108
<212> DNA

<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 518
ggtagctcta ccccttctgg tgcaaccggc tctccaggtg catccccggg taccagctct 60

accggttctc caggtactcc gggtagcggt accgcttctt cctctcca 108

<210> 519
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 519
Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
1 5 10 15

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
20 25 30

Thr Gly Ser Pro
35

<210> 520
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 520
ggtagctcta ctccgtctgg tgcaaccggt tccccaggta gctctacccc ttctggtgca 60

accggctccc caggtagctc tacccttct ggtgcaactg gctctcca 108

<210> 521
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 521
Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
1 5 10 15

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
20 25 30

Thr Gly Ser Pro

35

<210> 522
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 522
ggtgcttctc ctggcactag ctctaccggt tctccaggta cccctggtag cggtaccgca 60

tcttcctctc caggtagctc tactccttct ggtgctactg gttcccca 108

<210> 523
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 523
Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
1 5 10 15

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
20 25 30

Gly Thr Gly Pro
35

<210> 524
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 524
ggtagctcta ccccgtctgg tgctactggc tccccaggtt ctagcccttc tgcatccacc 60

ggtaccggtc caggttctag cccgtctgca tctactggta ctggtcca 108

<210> 525
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 525
Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
1 5 10 15

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
20 25 30

Gly Thr Gly Pro
35

<210> 526
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 526
ggtgcttccc cgggcaccag ctctactggt tctccaggtt ctagcccgtc tgcttctact 60

ggtactggtc caggttctag cccttctgct tccactggta ctggtcca 108

<210> 527
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 527
Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
1 5 10 15

Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala
20 25 30

Ser Ser Ser Pro
35

<210> 528
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 528
ggtgcttccc cgggtaccag ctctaccggt tctccaggtt ctagcccttc tgcttctacc 60

ggtaccggtc caggtacccc tggcagcggt accgcatctt cctctcca 108

<210> 529
<211> 432
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic

polynucleotide

<400> 529
ggtaccccgg gcagcggtac cgcatcttcc tctccaggta gctctacccc gtctggtgct 60

accggttccc caggtagctc taccccgtct ggtgcaaccg gctccccagg tagcccggct 120

ggctctccta cctctactga ggaaggtact tctgaaagcg ctactcctga gtctggtcca 180

ggtacctcta ctgaaccgtc cgaaggtagc gctccaggtt ctagcccttc tgcatccacc 240

ggtaccggcc caggttctag cccgtctgct tctaccggta ctggtccagg tgcttctccg 300

ggtactagct ctactggttc tccaggtacc tctaccgaac cgtccgaggg tagcgcacca 360

ggtacctcta ctgaaccgtc tgagggtagc gctccaggta gcgaaccggc aacctccggt 420

tctgaaactc ca 432

<210> 530
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 530
Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
1 5 10 15

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
20 25 30

Thr Gly Ser Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
35 40 45

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
50 55 60

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Ser Pro Ser Ala Ser Thr
65 70 75 80

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
85 90 95

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Ser Thr
100 105 110

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
115 120 125

Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
130 135 140

<210> 531
<211> 432

<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 531
ggttctacca gcgaatcccc ttctggcact gcaccaggtt ctactagcga atccccttct 60

ggtaccgcac caggtacttc tccgagcggc gaatcttcta ctgctccagg tacctctact 120

gaaccttccg aaggcagcgc tccaggtacc tctaccgaac cgtccgaggg cagcgcacca 180

ggtacttctg aaagcgcaac ccctgaatcc ggtccaggtg catctcctgg taccagctct 240

accggttctc caggtagctc tactccttct ggtgctactg gctctccagg tgcttccccg 300

ggtaccagct ctaccggttc tccaggttct actagcgaat ctccttctgg cactgcacca 360

ggttctacca gcgaatctcc gtctggcact gcaccaggta cctctacccc tgaaagcggt 420

tccgcttctc ca 432

<210> 532
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 532
Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser
1 5 10 15

Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser
20 25 30

Ser Thr Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
35 40 45

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
50 55 60

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
65 70 75 80

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
85 90 95

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Thr Ser
100 105 110

Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser
115 120 125

Gly Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro

130 135 140

<210> 533
<211> 432
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 533
ggtacttcta ccgaaccttc cgagggcagc gcaccaggta cttctgaaag cgctacccct 60

gagtccggcc caggtacttc tgaaagcgct actcctgaat ccggtccagg tacctctact 120

gaaccttctg agggcagcgc tccaggtact tctgaaagcg ctaccccgga gtccggtcca 180

ggtacttcta ctgaaccgtc cgaaggtagc gcaccaggta cttctactga accttccgaa 240

ggtagcgctc caggtagcga acctgctact tctggttctg aaaccccagg tagcccggct 300

ggctctccga cctccaccga ggaaggtgct tctcctggca ccagctctac tggttctcca 360

ggttctagcc cttctgcttc taccggtact ggtccaggtt ctagcccttc tgcatccact 420

ggtactggtc ca 432

<210> 534
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 534
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
20 25 30

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
35 40 45

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
50 55 60

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
65 70 75 80

Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
85 90 95

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ala Ser Pro
100 105 110

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
115 120 125

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
130 135 140

<210> 535
<211> 432
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 535
ggtagcgaac cggcaacctc tggctctgaa accccaggta cctctgaaag cgctactccg 60

gaatctggtc caggtacttc tgaaagcgct actccggaat ccggtccagg ttctaccagc 120

gaatctcctt ctggcaccgc tccaggttct actagcgaat ccccgtctgg taccgcacca 180

ggtacttctc ctagcggcga atcttctacc gcaccaggtg catctccggg tactagctct 240

accggttctc caggttctag cccttctgct tccactggta ccggcccagg tagctctacc 300

ccgtctggtg ctactggttc cccaggtagc tctactccgt ctggtgcaac cggttcccca 360

ggtagctcta ctccttctgg tgctactggc tccccaggtg catcccctgg caccagctct 420

accggttctc ca 432

<210> 536
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 536
Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
20 25 30

Glu Ser Gly Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
35 40 45

Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Pro
50 55 60

Ser Gly Glu Ser Ser Thr Ala Pro Gly Ala Ser Pro Gly Thr Ser Ser
65 70 75 80

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
85 90 95

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
100 105 110

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
115 120 125

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
130 135 140

<210> 537
<211> 432
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 537
ggtgcttctc cgggcaccag ctctactggt tctccaggtt ctagcccttc tgcatccacc 60

ggtaccggtc caggtagctc taccccttct ggtgcaaccg gctctccagg tacttctgaa 120

agcgctaccc cggaatctgg cccaggtagc gaaccggcta cttctggttc tgaaacccca 180

ggtagcgaac cggctacctc cggttctgaa actccaggta cttctgaaag cgctactccg 240

gagtccggtc caggtacctc taccgaaccg tccgaaggca gcgctccagg tacttctact 300

gaaccttctg agggtagcgc tccaggtacc tctaccgaac cgtccgaggg tagcgcacca 360

ggtacctcta ctgaaccgtc tgagggtagc gctccaggta gcgaaccggc aacctccggt 420

tctgaaactc ca 432

<210> 538
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 538
Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
1 5 10 15

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala
20 25 30

Thr Gly Ser Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
35 40 45

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
50 55 60

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
65 70 75 80

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
85 90 95

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
100 105 110

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
115 120 125

Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
130 135 140

<210> 539
<211> 432
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 539
ggtacctcta ccgaaccttc cgaaggtagc gctccaggta gcccggcagg ttctcctact 60

tccactgagg aaggtacttc taccgaacct tctgagggta gcgcaccagg tacctctgaa 120

agcgcaactc ctgagtctgg cccaggtagc gaacctgcta cctccggctc tgagactcca 180

ggtacctctg aaagcgcaac cccggaatct ggtccaggta gcccggctgg ctctcctacc 240

tctactgagg aaggtacttc tgaaagcgct actcctgagt ctggtccagg tacctctact 300

gaaccgtccg aaggtagcgc tccaggtagc gaacctgcta cttctggttc tgaaactcca 360

ggtacttcta ccgaaccgtc cgagggtagc gctccaggta gcgaacctgc tacttctggt 420

tctgaaactc ca 432

<210> 540
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 540
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
1 5 10 15

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
35 40 45

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu

50 55 60

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
65 70 75 80

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
85 90 95

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
100 105 110

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu
115 120 125

Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
130 135 140

<210> 541
<211> 432
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 541
ggtacttcta ccgaaccgtc cgaaggcagc gctccaggta cctctactga accttccgag 60

ggcagcgctc caggtacctc taccgaacct tctgaaggta gcgcaccagg tacttctacc 120

gaaccgtccg aaggcagcgc tccaggtacc tctactgaac cttccgaggg cagcgctcca 180

ggtacctcta ccgaaccttc tgaaggtagc gcaccaggta cttctaccga accttccgag 240

ggcagcgcac caggtacttc tgaaagcgct acccctgagt ccggcccagg tacttctgaa 300

agcgctactc ctgaatccgg tccaggtact tctactgaac cttccgaagg tagcgctcca 360

ggtagcgaac ctgctacttc tggttctgaa accccaggta gcccggctgg ctctccgacc 420

tccaccgagg aa 432

<210> 542
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 542
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
1 5 10 15

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
35 40 45

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
50 55 60

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
65 70 75 80

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
85 90 95

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
100 105 110

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
115 120 125

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
130 135 140

<210> 543
<211> 432
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 543
ggtacttcta ccgaaccgtc cgagggcagc gctccaggta cttctactga accttctgaa 60

ggcagcgctc caggtacttc tactgaacct tccgaaggta gcgcaccagg ttctaccagc 120

gaatcccctt ctggtactgc tccaggttct accagcgaat ccccttctgg caccgcacca 180

ggtacttcta cccctgaaag cggctccgct tctccaggta gcgaacctgc aacctctggc 240

tctgaaaccc caggtacctc tgaaagcgct actcctgaat ctggcccagg tacttctact 300

gaaccgtccg agggcagcgc accaggtact tctactgaac cgtctgaagg tagcgcacca 360

ggtacttctg aaagcgcaac cccggaatcc ggcccaggta cctctgaaag cgcaaccccg 420

gagtccggcc ca 432

<210> 544
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 544
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
1 5 10 15

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
35 40 45

Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Thr
50 55 60

Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Glu Pro Ala Thr Ser Gly
65 70 75 80

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
85 90 95

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
100 105 110

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
115 120 125

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
130 135 140

<210> 545
<211> 432
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 545
ggtagctcta ccccttctgg tgctaccggc tctccaggtt ctagcccgtc tgcttctacc 60

ggtaccggtc caggtagctc tacccccttct ggtgctactg gttctccagg tagccctgct 120

ggctctccga cttctactga ggaaggtagc ccggctggtt ctccgacttc tactgaggaa 180

ggtacttcta ccgaaccttc cgaaggtagc gctccaggtg cttccccggg cactagctct 240

accggttctc caggttctag cccttctgca tctactggta ctggcccagg tactccgggc 300

agcggtactg cttcttcctc tccaggttct actagctcta ctgctgaatc tcctggccca 360

ggtacttctc ctagcggtga atcttctacc gctccaggta cctctactcc ggaaagcggt 420

tctgcatctc ca 432

<210> 546
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 546
Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
1   5   10   15

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala
            20   25   30

Thr Gly Ser Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
        35   40   45

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
    50   55   60

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Pro Gly Thr Ser Ser
65   70   75   80

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
                85   90   95

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Thr Ser
            100   105   110

Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro Ser Gly Glu Ser
        115   120   125

Ser Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
    130   135   140

<210> 547
<211> 432
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
      polynucleotide

<400> 547
ggtacctcta ctgaaccttc tgagggcagc gctccaggta cttctgaaag cgctaccccg 60

gagtccggtc caggtacttc tactgaaccg tccgaaggta gcgcaccagg tacttctact 120

gaaccgtctg aaggtagcgc accaggtact tctgaaagcg caaccccgga atccggccca 180

ggtacctctg aaagcgcaac cccggagtcc ggcccaggta ctcctggcag cggtaccgct 240

tcttcttctc caggtgcttc tcctggtact agctctactg gttctccagg tgcttctccg 300

ggcactagct ctactggttc tccaggtagc cctgctggct ctccgacttc tactgaggaa 360

ggtagcccgg ctggttctcc gacttctact gaggaaggta cttctaccga accttccgaa 420

ggtagcgctc ca   432

<210> 548
<211> 144
<212> PRT

<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 548
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
35 40 45

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu
50 55 60

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala
65 70 75 80

Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
85 90 95

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Pro Ala
100 105 110

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
115 120 125

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
130 135 140

<210> 549
<211> 432
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 549
ggtagcccag caggctctcc gacttccact gaggaaggta cttctactga accttccgaa 60

ggcagcgcac caggtacctc tactgaacct tctgagggca gcgctccagg tacctctacc 120

gaaccgtctg aaggtagcgc accaggtacc tctgaaagcg caactcctga gtccggtcca 180

ggtacttctg aaagcgcaac cccggagtct ggcccaggta ccccgggtag cggtactgct 240

tcttcctctc caggtagctc tacccttct ggtgcaaccg gctctccagg tgcttctccg 300

ggcaccagct ctaccggttc tccaggtacc tctactgaac cttctgaggg cagcgctcca 360

ggtacttctg aaagcgctac cccggagtcc ggtccaggta cttctactga accgtccgaa 420

ggtagcgcac ca 432

<210> 550
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 550
Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
1 5 10 15

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
35 40 45

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu
50 55 60

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala
65 70 75 80

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
85 90 95

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Ser Thr
100 105 110

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
115 120 125

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
130 135 140

<210> 551
<211> 432
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 551
ggtagcgaac cggcaacctc cggctctgaa actccaggta cttctgaaag cgctactccg 60

gaatccggcc caggtagcga accggctact tccggctctg aaaccccagg tagctctacc 120

ccgtctggtg caaccggctc cccaggtact cctggtagcg gtaccgcttc ttcttctcca 180

ggtagctcta ctccgtctgg tgctaccggc tccccaggtg catctcctgg taccagctct 240

accggttctc caggtagctc tactccttct ggtgctactg gctctccagg tgcttccccg 300

ggtaccagct ctaccggttc tccaggtagc gaacctgcta cttctggttc tgaaactcca 360

ggtacttcta ccgaaccgtc cgagggtagc gctccaggta gcgaacctgc tacttctggt 420

tctgaaactc ca 432

<210> 552
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 552
Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
20 25 30

Ser Glu Thr Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
35 40 45

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
50 55 60

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
65 70 75 80

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
85 90 95

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Glu Pro
100 105 110

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu
115 120 125

Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
130 135 140

<210> 553
<211> 432
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 553
ggtacctcta ctgaaccttc cgaaggcagc gctccaggta cctctaccga accgtccgag 60

ggcagcgcac caggtacttc tgaaagcgca acccctgaat ccggtccagg tagccctgct 120

ggctctccga cttctactga ggaaggtagc ccggctggtt ctccgacttc tactgaggaa 180

ggtacttcta ccgaaccttc cgaaggtagc gctccaggta gcccggctgg ttctccgact 240

tccaccgagg aaggtacctc tactgaacct tctgagggta gcgctccagg tacctctact 300

gaaccttccg aaggcagcgc tccaggtgct tccccgggca ccagctctac tggttctcca 360

ggttctagcc cgtctgcttc tactggtact ggtccaggtt ctagcccttc tgcttccact 420

ggtactggtc ca 432

<210> 554
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 554
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
1 5 10 15

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
20 25 30

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
35 40 45

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
50 55 60

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
65 70 75 80

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
85 90 95

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Pro
100 105 110

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
115 120 125

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
130 135 140

<210> 555
<211> 432
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 555

ggtagctcta ccccgtctgg tgctaccggt tccccaggtg cttctcctgg tactagctct 60

accggttctc caggtagctc taccccgtct ggtgctactg gctctccagg tagccctgct 120

ggctctccaa cctccaccga agaaggtacc tctgaaagcg caacccctga atccggccca 180

ggtagcgaac cggcaacctc cggttctgaa accccaggtg catctcctgg tactagctct 240

actggttctc caggtagctc tactccgtct ggtgcaaccg gctctccagg ttctagccct 300

tctgcatcta ccggtactgg tccaggttct accagcgaat ccccttctgg tactgctcca 360

ggttctacca gcgaatcccc ttctggcacc gcaccaggta cttctacccc tgaaagcggc 420

tccgcttctc ca 432

<210> 556
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 556

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro
1 5 10 15

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
20 25 30

Thr Gly Ser Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
35 40 45

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
50 55 60

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ala Ser Pro Gly Thr Ser Ser
65 70 75 80

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
85 90 95

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Thr Ser
100 105 110

Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser
115 120 125

Gly Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
130 135 140

<210> 557
<211> 432
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 557
ggttctacca gcgaatctcc ttctggcacc gctccaggtt ctactagcga atccccgtct 60

ggtaccgcac caggtacttc tcctagcggc gaatcttcta ccgcaccagg tacctctgaa 120

agcgctactc cggagtctgg cccaggtacc tctactgaac cgtctgaggg tagcgctcca 180

ggtacttcta ctgaaccgtc cgaaggtagc gcaccaggta cctctactga accttctgag 240

ggcagcgctc caggtacttc tgaaagcgct accccggagt ccggtccagg tacttctact 300

gaaccgtccg aaggtagcgc accaggtagc tctaccccgt ctggtgctac cggttcccca 360

ggtgcttctc ctggtactag ctctaccggt tctccaggta gctctacccc gtctggtgct 420

actggctctc ca 432

<210> 558
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 558
Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser
1 5 10 15

Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser
20 25 30

Ser Thr Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
35 40 45

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
50 55 60

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
65 70 75 80

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
85 90 95

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Ser Thr
100 105 110

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
115 120 125

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
130 135 140

<210> 559
<211> 432
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 559
ggttctacta gctctactgc agaatctccg ggcccaggta cctctcctag cggtgaatct 60

tctaccgctc caggtacttc tccgagcggt gaatcttcta ccgctccagg ttctactagc 120

tctaccgctg aatctccggg tccaggttct accagctcta ctgcagaatc tcctggccca 180

ggtacttcta ctccggaaag cggttccgct tctccaggta cttctcctag cggtgaatct 240

tctaccgctc caggttctac cagctctact gctgaatctc ctggcccagg tacttctacc 300

ccggaaagcg gctccgcttc tccaggttct accagctcta ccgctgaatc tcctggccca 360

ggttctacta gcgaatctcc gtctggcacc gcaccaggta cttcccctag cggtgaatct 420

tctactgcac ca 432

<210> 560
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 560
Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro
1 5 10 15

Ser Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser
20 25 30

Ser Thr Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro
35 40 45

Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Thr
50 55 60

Pro Glu Ser Gly Ser Ala Ser Pro Gly Thr Ser Pro Ser Gly Glu Ser
65 70 75 80

Ser Thr Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro
85 90 95

Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser
100 105 110

Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser Glu Ser Pro Ser
115 120 125

Gly Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro
130 135 140

<210> 561
<211> 432
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 561
ggtacctcta ctccggaaag cggttccgca tctccaggtt ctaccagcga atccccgtct 60

ggcaccgcac caggttctac tagctctact gctgaatctc cgggcccagg tacctctact 120

gaaccttccg aaggcagcgc tccaggtacc tctaccgaac cgtccgaggg cagcgcacca 180

ggtacttctg aaagcgcaac ccctgaatcc ggtccaggta cctctgaaag cgctactccg 240

gagtctggcc caggtacctc tactgaaccg tctgagggta gcgctccagg tacttctact 300

gaaccgtccg aaggtagcgc accaggtact tctgaaagcg ctactccgga gtccggtcca 360

ggtacctcta ccgaaccgtc cgaaggcagc gctccaggta cttctactga accttctgag 420

ggtagcgctc cc 432

<210> 562
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 562
Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser
1 5 10 15

Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu
20 25 30

Ser Pro Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
35 40 45

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
50 55 60

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
65 70 75 80

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
85 90 95

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
100 105 110

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
115 120 125

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
130 135 140

<210> 563
<211> 432
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 563
ggtacctcta ccgaaccgtc cgagggtagc gcaccaggta cctctactga accgtctgag 60

ggtagcgctc caggtagcga accggcaacc tccggttctg aaactccagg tacttctact 120

gaaccgtctg aaggtagcgc accaggtact tctgaaagcg caaccccgga atccggccca 180

ggtacctctg aaagcgcaac cccggagtcc ggcccaggtg catctccggg tactagctct 240

accggttctc caggttctag cccttctgct tccactggta ccggcccagg tagctctacc 300

ccgtctggtg ctactggttc cccaggtagc tctactccgt ctggtgcaac cggttcccca 360

ggtagctcta ctccttctgg tgctactggc tccccaggtg catcccctgg caccagctct 420

accggttctc ca 432

<210> 564
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 564
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
1 5 10 15

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
20 25 30

Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
35 40 45

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu
50 55 60

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
65 70 75 80

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro

85 90 95

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
100 105 110

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
115 120 125

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
130 135 140

<210> 565
<211> 432
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 565
ggtagcgaac cggcaacctc tggctctgaa actccaggta gcgaacctgc aacctccggc 60

tctgaaaccc caggtacttc tactgaacct tctgagggca gcgcaccagg tagcgaacct 120

gcaacctctg gctctgaaac cccaggtacc tctgaaagcg ctactcctga atctggccca 180

ggtacttcta ctgaaccgtc cgagggcagc gcaccaggta gctctactcc gtctggtgct 240

accggctctc caggtagctc tacccttct ggtgcaaccg gctccccagg tgcttctccg 300

ggtaccagct ctactggttc tccaggtagc tctaccccgt ctggtgctac cggttcccca 360

ggtgcttctc ctggtactag ctctaccggt tctccaggta gctctacccc gtctggtgct 420

actggctctc ca 432

<210> 566
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 566
Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
1 5 10 15

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
35 40 45

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
50 55 60

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Ser Thr Pro Ser Gly Ala
65 70 75 80

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
85 90 95

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
100 105 110

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
115 120 125

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
130 135 140

<210> 567
<211> 432
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 567
ggtacttcta ccgaaccgtc cgagggcagc gctccaggta cttctactga accttctgaa 60

ggcagcgctc caggtacttc tactgaacct tccgaaggta gcgcaccagg tacttctgaa 120

agcgctactc cggagtccgg tccaggtacc tctaccgaac cgtccgaagg cagcgctcca 180

ggtacttcta ctgaaccttc tgagggtagc gctccaggtt ctactagcga atctccgtct 240

ggcactgctc caggtacttc tcctagcggt gaatcttcta ccgctccagg tacttcccct 300

agcggcgaat cttctaccgc tccaggtagc ccggctggct ctcctacctc tactgaggaa 360

ggtacttctg aaagcgctac tcctgagtct ggtccaggta cctctactga accgtccgaa 420

ggtagcgctc ca 432

<210> 568
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 568
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
1 5 10 15

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
35 40 45

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
50 55 60

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser
65 70 75 80

Gly Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro
85 90 95

Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Pro Ala
100 105 110

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
115 120 125

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
130 135 140

<210> 569
<211> 432
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 569
ggtacttcta ctgaaccttc cgaaggtagc gctccaggta gcgaacctgc tacttctggt 60

tctgaaaccc caggtagccc ggctggctct ccgacctcca ccgaggaagg tagcccggca 120

ggctctccga cctctactga ggaaggtact tctgaaagcg caaccccgga gtccggccca 180

ggtacctcta ccgaaccgtc tgagggcagc gcaccaggta cctctactga accttccgaa 240

ggcagcgctc caggtacctc taccgaaccg tccgagggca gcgcaccagg tacttctgaa 300

agcgcaaccc ctgaatccgg tccaggtagc tctactccgt ctggtgcaac cggctcccca 360

ggttctagcc cgtctgcttc cactggtact ggcccaggtg cttccccggg caccagctct 420

actggttctc ca 432

<210> 570
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 570
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
1 5 10 15

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
20 25 30

Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
35 40 45

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
50 55 60

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
65 70 75 80

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
85 90 95

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Ser Thr
100 105 110

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
115 120 125

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
130 135 140

<210> 571
<211> 432
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 571
ggtagcgaac cggctacttc cggctctgag actccaggta gccctgctgg ctctccgacc 60

tctaccgaag aaggtacctc tgaaagcgct acccctgagt ctggcccagg tacctctact 120

gaaccttccg aaggcagcgc tccaggtacc tctaccgaac cgtccgaggg cagcgcacca 180

ggtacttctg aaagcgcaac ccctgaatcc ggtccaggta cctctactcc ggaaagcggt 240

tccgcatctc caggttctac cagcgaatcc ccgtctggca ccgcaccagg ttctactagc 300

tctactgctg aatctccggg cccaggtact tctgaaagcg ctactccgga gtccggtcca 360

ggtacctcta ccgaaccgtc cgaaggcagc gctccaggta cttctactga accttctgag 420

ggtagcgctc ca 432

<210> 572
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 572
Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala

1 5 10 15

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
20 25 30

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
35 40 45

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
50 55 60

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Pro Glu Ser Gly
65 70 75 80

Ser Ala Ser Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
85 90 95

Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Glu
100 105 110

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
115 120 125

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
130 135 140

<210> 573
<211> 432
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 573
ggtacttcta ccgaaccgtc cgagggcagc gctccaggta cttctactga accttctgaa 60

ggcagcgctc caggtacttc tactgaacct tccgaaggta gcgcaccagg tacctctacc 120

gaaccgtctg aaggtagcgc accaggtacc tctgaaagcg caactcctga gtccggtcca 180

ggtacttctg aaagcgcaac cccggagtct ggcccaggta ctcctggcag cggtaccgca 240

tcttcctctc caggtagctc tactccgtct ggtgcaactg gttccccagg tgcttctccg 300

ggtaccagct ctaccggttc tccaggttcc accagctcta ctgctgaatc tcctggtcca 360

ggtacctctc ctagcggtga atcttctact gctccaggta cttctactcc tgaaagcggc 420

tctgcttctc ca 432

<210> 574
<211> 144
<212> PRT
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 574

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
1 5 10 15

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
35 40 45

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu
50 55 60

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala
65 70 75 80

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
85 90 95

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Thr Ser
100 105 110

Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro Ser Gly Glu Ser
115 120 125

Ser Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
130 135 140

<210> 575
<211> 432
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 575

ggtagcccgg caggctctcc gacctctact gaggaaggta cttctgaaag cgcaaccccg 60

gagtccggcc caggtacctc taccgaaccg tctgagggca gcgcaccagg tacttctgaa 120

agcgcaaccc ctgaatccgg tccaggtagc gaaccggcta cttctggctc tgagactcca 180

ggtacttcta ccgaaccgtc cgaaggtagc gcaccaggta gcccggctgg ttctccgact 240

tccaccgagg aaggtacctc tactgaacct tctgagggta gcgctccagg tacctctact 300

gaaccttccg aaggcagcgc tccaggtact tctaccgaac cgtccgaggg cagcgctcca 360

ggtacttcta ctgaaccttc tgaaggcagc gctccaggta cttctactga accttccgaa 420

ggtagcgcac ca 432

<210> 576

<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 576
Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
35 40 45

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr
50 55 60

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
65 70 75 80

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
85 90 95

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
100 105 110

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
115 120 125

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
130 135 140

<210> 577
<211> 432
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 577
ggtacttctc cgagcggtga atcttctacc gcaccaggtt ctactagctc taccgctgaa 60

tctccgggcc caggtacttc tccgagcggt gaatcttcta ctgctccagg tacctctgaa 120

agcgctactc cggagtctgg cccaggtacc tctactgaac cgtctgaggg tagcgctcca 180

ggtacttcta ctgaaccgtc cgaaggtagc gcaccaggtt ctagcccttc tgcatctact 240

ggtactggcc caggtagctc tactccttct ggtgctaccg gctctccagg tgcttctccg 300

ggtactagct ctaccggttc tccaggtact tctactccgg aaagcggttc cgcatctcca 360

ggtacttctc ctagcggtga atcttctact gctccaggta cctctcctag cggcgaatct 420

tctactgctc ca 432

<210> 578
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 578
Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser
1 5 10 15

Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro Ser Gly Glu Ser
20 25 30

Ser Thr Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
35 40 45

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
50 55 60

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Ser Pro Ser Ala Ser Thr
65 70 75 80

Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
85 90 95

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Ser Thr
100 105 110

Pro Glu Ser Gly Ser Ala Ser Pro Gly Thr Ser Pro Ser Gly Glu Ser
115 120 125

Ser Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro
130 135 140

<210> 579
<211> 432
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 579
ggtacctctg aaagcgctac tccggagtct ggcccaggta cctctactga accgtctgag 60

ggtagcgctc caggtacttc tactgaaccg tccgaaggta gcgcaccagg tagccctgct 120

ggctctccga cttctactga ggaaggtagc ccggctggtt ctccgacttc tactgaggaa 180

ggtacttcta ccgaaccttc cgaaggtagc gctccaggtt ctagcccttc tgcttccacc 240

ggtactggcc caggtagctc taccccttct ggtgctaccg gctccccagg tagctctact 300

ccttctggtg caactggctc tccaggtagc gaaccggcaa cttccggctc tgaaacccca 360

ggtacttctg aaagcgctac tcctgagtct ggcccaggta gcgaacctgc tacctctggc 420

tctgaaaccc ca 432

<210> 580
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 580

Gly	Thr	Ser	Glu	Ser	Ala	Thr	Pro	Glu	Ser	Gly	Pro	Gly	Thr	Ser	Thr
1				5				10					15		
Glu	Pro	Ser	Glu	Gly	Ser	Ala	Pro	Gly	Thr	Ser	Thr	Glu	Pro	Ser	Glu
			20					25					30		
Gly	Ser	Ala	Pro	Gly	Ser	Pro	Ala	Gly	Ser	Pro	Thr	Ser	Thr	Glu	Glu
		35					40					45			
Gly	Ser	Pro	Ala	Gly	Ser	Pro	Thr	Ser	Thr	Glu	Glu	Gly	Thr	Ser	Thr
	50					55					60				
Glu	Pro	Ser	Glu	Gly	Ser	Ala	Pro	Gly	Ser	Ser	Pro	Ser	Ala	Ser	Thr
65					70					75					80
Gly	Thr	Gly	Pro	Gly	Ser	Ser	Thr	Pro	Ser	Gly	Ala	Thr	Gly	Ser	Pro
				85					90					95	
Gly	Ser	Ser	Thr	Pro	Ser	Gly	Ala	Thr	Gly	Ser	Pro	Gly	Ser	Glu	Pro
			100					105					110		
Ala	Thr	Ser	Gly	Ser	Glu	Thr	Pro	Gly	Thr	Ser	Glu	Ser	Ala	Thr	Pro
		115					120					125			
Glu	Ser	Gly	Pro	Gly	Ser	Glu	Pro	Ala	Thr	Ser	Gly	Ser	Glu	Thr	Pro
	130					135					140				

<210> 581
<211> 432
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 581
ggtacttcta ccgaaccgtc cgaaggcagc gctccaggta cctctactga accttccgag 60

ggcagcgctc caggtacctc taccgaacct tctgaaggta gcgcaccagg tagctctacc 120

ccgtctggtg ctaccggttc cccaggtgct tctcctggta ctagctctac cggttctcca 180

ggtagctcta ccccgtctgg tgctactggc tctccaggta cttctgaaag cgcaacccct 240

gaatccggtc caggtagcga accggctact tctggctctg agactccagg tacttctacc 300

gaaccgtccg aaggtagcgc accaggttct actagcgaat ctccttctgg cactgcacca 360

ggttctacca gcgaatctcc gtctggcact gcaccaggta cctctacccc tgaaagcggt 420

tccgcttctc ca 432

<210> 582
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 582
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
1 5 10 15

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
35 40 45

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
50 55 60

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Ser Glu Ser Ala Thr Pro
65 70 75 80

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
85 90 95

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Thr Ser
100 105 110

Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser
115 120 125

Gly Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
130 135 140

<210> 583
<211> 432
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 583
ggtacctcta ctcctgaaag cggttctgca tctccaggtt ccactagctc taccgcagaa 60

tctccgggcc caggttctac tagctctact gctgaatctc ctggcccagg ttctagccct 120

tctgcatcta ctggtactgg cccaggtagc tctactcctt ctggtgctac cggctctcca 180

ggtgcttctc cgggtactag ctctaccggt tctccaggta gcgaaccggc aacctccggc 240

tctgaaaccc caggtacctc tgaaagcgct actcctgaat ccggcccagg tagcccggca 300

ggttctccga cttccactga ggaaggttct actagcgaat ctccttctgg cactgcacca 360

ggttctacca gcgaatctcc gtctggcact gcaccaggta cctctacccc tgaaagcggt 420

tccgcttctc cc 432

<210> 584
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 584
Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser
1 5 10 15

Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser Ser Thr Ala Glu
20 25 30

Ser Pro Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
35 40 45

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro
50 55 60

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Glu Pro Ala Thr Ser Gly
65 70 75 80

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
85 90 95

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Thr Ser
100 105 110

Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser
115 120 125

Gly Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
130 135 140

<210> 585
<211> 432
<212> DNA

<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 585
ggtagcccgg ctggctctcc tacctctact gaggaaggta cttctgaaag cgctactcct 60

gagtctggtc caggtacctc tactgaaccg tccgaaggta gcgctccagg ttctaccagc 120

gaatctcctt ctggcaccgc tccaggttct actagcgaat ccccgtctgg taccgcacca 180

ggtacttctc ctagcggcga atcttctacc gcaccaggta cctctaccga accttccgaa 240

ggtagcgctc caggtagccc ggcaggttct cctacttcca ctgaggaagg tacttctacc 300

gaaccttctg agggtagcgc accaggtagc gaacctgcaa cctctggctc tgaaacccca 360

ggtacctctg aaagcgctac tcctgaatct ggcccaggta cttctactga accgtccgag 420

ggcagcgcac ca 432

<210> 586
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 586
Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
35 40 45

Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Pro
50 55 60

Ser Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
65 70 75 80

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
85 90 95

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
100 105 110

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
115 120 125

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
130 135 140

<210> 587
<211> 432
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 587
ggtacctcta ccgaaccttc cgaaggtagc gctccaggta gcccggcagg ttctcctact 60

tccactgagg aaggtacttc taccgaacct tctgagggta gcgcaccagg tacctcccct 120

agcggcgaat cttctactgc tccaggtacc tctcctagcg gcgaatcttc taccgctcca 180

ggtacctccc ctagcggtga atcttctacc gcaccaggtt ctaccagcga atcccttct 240

ggtactgctc caggttctac cagcgaatcc ccttctggca ccgcaccagg tacttctacc 300

cctgaaagcg gctccgcttc tccaggtagc gaacctgcaa cctctggctc tgaaacccca 360

ggtacctctg aaagcgctac tcctgaatct ggcccaggta cttctactga accgtccgag 420

ggcagcgcac ca 432

<210> 588
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 588
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
1 5 10 15

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro
35 40 45

Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Pro
50 55 60

Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser
65 70 75 80

Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
85 90 95

Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Glu Pro
100 105 110

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro

115 120 125

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
130 135 140

<210> 589
<211> 432
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 589
ggtagcgaac cggcaacctc tggctctgaa accccaggta cctctgaaag cgctactccg 60

gaatctggtc caggtacttc tgaaagcgct actccggaat ccggtccagg tacttctccg 120

agcggtgaat cttctaccgc accaggttct actagctcta ccgctgaatc tccgggccca 180

ggtacttctc cgagcggtga atcttctact gctccaggtt ctactagcga atccccgtct 240

ggtactgctc caggtacttc ccctagcggt gaatcttcta ctgctccagg ttctaccagc 300

tctaccgcag aatctccggg tccaggtagc tctactccgt ctggtgcaac cggttcccca 360

ggtagctcta ccccttctgg tgcaaccggc tccccaggta gctctacccc ttctggtgca 420

aactggctct cc 432

<210> 590
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 590
Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
20 25 30

Glu Ser Gly Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro
35 40 45

Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro
50 55 60

Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser
65 70 75 80

Gly Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro
85 90 95

Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Ser Thr
100 105 110

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
115 120 125

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Asn Trp Leu Ser
130 135 140

<210> 591
<211> 432
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 591
ggtagccctg ctggctctcc gacttctact gaggaaggta gcccggctgg ttctccgact 60

tctactgagg aaggtacttc taccgaacct tccgaaggta gcgctccagg tacctctact 120

gaaccttccg aaggcagcgc tccaggtacc tctaccgaac cgtccgaggg cagcgcacca 180

ggtacttctg aaagcgcaac ccctgaatcc ggtccaggtg catctcctgg taccagctct 240

accggttctc caggtagctc tactccttct ggtgctactg gctctccagg tgcttccccg 300

ggtaccagct ctaccggttc tccaggtagc tctaccccgt ctggtgctac tggttctcca 360

ggtactccgg gcagcggtac tgcttcttcc tctccaggta gctctacccc ttctggtgct 420

actggctctc ca 432

<210> 592
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 592
Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala
1 5 10 15

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
35 40 45

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
50 55 60

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
65 70 75 80

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
85 90 95

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
100 105 110

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
115 120 125

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
130 135 140

<210> 593
<211> 432
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 593
ggtagctcta ccccgtctgg tgctactggt tctccaggta ctccgggcag cggtactgct 60

tcttcctctc caggtagctc tacccccttct ggtgctactg gctctccagg tagcccggct 120

ggctctccta cctctactga ggaaggtact tctgaaagcg ctactcctga gtctggtcca 180

ggtacctcta ctgaaccgtc cgaaggtagc gctccaggta cctctgaaag cgcaactcct 240

gagtctggcc caggtagcga acctgctacc tccggctctg agactccagg tacctctgaa 300

agcgcaaccc cggaatctgg tccaggtact tctactgaac cgtctgaagg tagcgcacca 360

ggtacttctg aaagcgcaac cccggaatcc ggcccaggta cctctgaaag cgcaaccccg 420

gagtccggcc ca 432

<210> 594
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 594
Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
1 5 10 15

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
20 25 30

Thr Gly Ser Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
35 40 45

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
50 55 60

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
65 70 75 80

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
85 90 95

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
100 105 110

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
115 120 125

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
130 135 140

<210> 595
<211> 1469
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 595
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

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Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415
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Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685
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Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960
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Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
               965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
           980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
       995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
   1010                1015                1020

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
   1025                1030                1035

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
   1040                1045                1050

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
   1055                1060                1065

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
   1070                1075                1080

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
   1085                1090                1095

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
   1100                1105                1110

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
   1115                1120                1125

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
   1130                1135                1140

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
   1145                1150                1155

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
   1160                1165                1170

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
   1175                1180                1185

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
   1190                1195                1200

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
   1205                1210                1215
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Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1220                 1225                 1230

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1235                 1240                 1245

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1250                 1255                 1260

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1265                 1270                 1275

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1280                 1285                 1290

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1295                 1300                 1305

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1310                 1315                 1320

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1325                 1330                 1335

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1340                 1345                 1350

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1355                 1360                 1365

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1370                 1375                 1380

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1385                 1390                 1395

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1400                 1405                 1410

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1415                 1420                 1425

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1430                 1435                 1440

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
    1445                 1450                 1455

Ser Pro  Ala Thr Gly His His  His His His His His
    1460                 1465
```

<210> 596
<211> 4410
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 596
gccaccagaa gatactacct gggtgcagtg gaactgtcat gggactatat gcaaagtgat 60

ctcggtgagc tgcctgtgga cgcaagattt cctcctagag tgccaaaatc ttttccattc 120

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<210> 597
<211> 1511
<212> PRT
<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic polypeptide

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Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Ala Ser Ser Arg Tyr Tyr
35 40 45

Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr Met Gln Ser Asp Leu Gly
50 55 60

Glu Leu Pro Val Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe
65 70 75 80

Pro Phe Asn Thr Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe
85 90 95

Thr Asp His Leu Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly
100 105 110

Leu Leu Gly Pro Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile
115 120 125

Thr Leu Lys Asn Met Ala Ser His Pro Val Ser Leu His Ala Val Gly
130 135 140

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Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr
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Ser Gln Arg Glu Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His
                165                 170                 175

Thr Tyr Val Trp Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp
            180                 185                 190

Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys
        195                 200                 205

Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly
    210                 215                 220

Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu
225                 230                 235                 240

Phe Ala Val Phe Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn
                245                 250                 255

Ser Leu Met Gln Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys
            260                 265                 270

Met His Thr Val Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile
        275                 280                 285

Gly Cys His Arg Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr
    290                 295                 300

Thr Pro Glu Val His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val
305                 310                 315                 320

Arg Asn His Arg Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu
                325                 330                 335

Thr Ala Gln Thr Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys
            340                 345                 350

His Ile Ser Ser His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val
        355                 360                 365

Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu
    370                 375                 380

Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val
385                 390                 395                 400

Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val
                405                 410                 415
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Ala Lys Lys His Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu
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Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser
        435                 440                 445

Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys
    450                 455                 460

Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr
465                 470                 475                 480

Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr
                485                 490                 495

Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser
            500                 505                 510

Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu
        515                 520                 525

Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro
    530                 535                 540

Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu
545                 550                 555                 560

Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser
                565                 570                 575

Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro
            580                 585                 590

Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile
        595                 600                 605

Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn
    610                 615                 620

Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro
625                 630                 635                 640

Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met
                645                 650                 655

His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys
            660                 665                 670

Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr
        675                 680                 685
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Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met
    690                 695                 700

Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val
705                 710                 715                 720

Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn
                725                 730                 735

Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser
            740                 745                 750

Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile
        755                 760                 765

Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe
    770                 775                 780

Ser Gln Asn Pro Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg
785                 790                 795                 800

Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
                805                 810                 815

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu
            820                 825                 830

Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile
        835                 840                 845

Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His
    850                 855                 860

Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys
865                 870                 875                 880

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
                885                 890                 895

Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
            900                 905                 910

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser
        915                 920                 925

Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln
    930                 935                 940

Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
945                 950                 955                 960
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Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp
                965                 970                 975

Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu
            980                 985                 990

Lys Asp Val His Ser Gly Leu Ile  Gly Pro Leu Leu Val  Cys His Thr
        995                 1000                1005

Asn Thr  Leu Asn Pro Ala His  Gly Arg Gln Val Thr  Val Gln Glu
    1010                1015                1020

Phe Ala  Leu Phe Phe Thr Ile  Phe Asp Glu Thr Lys  Ser Trp Tyr
    1025                1030                1035

Phe Thr  Glu Asn Met Glu Arg  Asn Cys Arg Ala Pro  Cys Asn Ile
    1040                1045                1050

Gln Met  Glu Asp Pro Thr Phe  Lys Glu Asn Tyr Arg  Phe His Ala
    1055                1060                1065

Ile Asn  Gly Tyr Ile Met Asp  Thr Leu Pro Gly Leu  Val Met Ala
    1070                1075                1080

Gln Asp  Gln Arg Ile Arg Trp  Tyr Leu Leu Ser Met  Gly Ser Asn
    1085                1090                1095

Glu Asn  Ile His Ser Ile His  Phe Ser Gly His Val  Phe Thr Val
    1100                1105                1110

Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
    1115                1120                1125

Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    1130                1135                1140

Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    1145                1150                1155

Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    1160                1165                1170

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
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Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
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Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
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Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
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Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    1235                1240                1245

Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
    1250                1255                1260

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
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Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
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Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
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Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
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Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
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Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
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Ser Lys  Ala Arg Leu His Leu  Gln Gly Arg Ser Asn  Ala Trp Arg
    1355                1360                1365

Pro Gln  Val Asn Asn Pro Lys  Glu Trp Leu Gln Val  Asp Phe Gln
    1370                1375                1380

Lys Thr  Met Lys Val Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser
    1385                1390                1395

Leu Leu  Thr Ser Met Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser
    1400                1405                1410

Gln Asp  Gly His Gln Trp Thr  Leu Phe Phe Gln Asn  Gly Lys Val
    1415                1420                1425

Lys Val  Phe Gln Gly Asn Gln  Asp Ser Phe Thr Pro  Val Val Asn
    1430                1435                1440

Ser Leu  Asp Pro Pro Leu Leu  Thr Arg Tyr Leu Arg  Ile His Pro
    1445                1450                1455

Gln Ser  Trp Val His Gln Ile  Ala Leu Arg Met Glu  Val Leu Gly
    1460                1465                1470
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Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu
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Thr Gly His His His His His His
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<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic polynucleotide

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agttgcagca tgccattggg aatggagagt aaagcaatat cagatgcaca gattactgct 4020

tcatcctact ttaccaatat gtttgccacc tggtctcctt caaaagctcg acttcacctc 4080

caagggagga gtaatgcctg gagacctcag gtgaataatc caaaagagtg gctgcaagtg 4140

gacttccaga agacaatgaa agtcacagga gtaactactc agggagtaaa atctctgctt 4200

accagcatgt atgtgaagga gttcctcatc tccagcagtc aagatggcca tcagtggact 4260

ctctttttc agaatggcaa agtaaaggtt tttcagggaa atcaagactc cttcacacct 4320

gtggtgaact ctctagaccc accgttactg actcgctacc ttcgaattca cccccagagt 4380

tgggtgcacc agattgccct gaggatggag gttctgggct gcgaggcaca ggacctctac 4440

ggcgccggat cacctggggc cgaaacggcc gaacaaaaac tcatctcaga agaggatctg 4500

tcacctgcaa ccggtcatca tcaccatcac cattga 4536

<210> 599
<211> 1514
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 599
Ala Thr Arg Gly Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr
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Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
20 25 30

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Ala Ser Ser Ala Thr Arg
35 40 45

Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr Met Gln Ser
50 55 60

Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro Arg Val Pro
65 70 75 80

```
Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys Thr Leu Phe
                85                  90                  95

Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro Arg Pro Pro
            100                 105                 110

Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val Tyr Asp Thr
        115                 120                 125

Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val Ser Leu His
    130                 135                 140

Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala Glu Tyr Asp
145                 150                 155                 160

Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val Phe Pro Gly
                165                 170                 175

Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn Gly Pro Met
            180                 185                 190

Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser His Val Asp
        195                 200                 205

Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu Leu Val Cys
    210                 215                 220

Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu His Lys Phe
225                 230                 235                 240

Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp His Ser Glu
                245                 250                 255

Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser Ala Arg Ala
            260                 265                 270

Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg Ser Leu Pro
        275                 280                 285

Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His Val Ile Gly
    290                 295                 300

Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu Gly His Thr
305                 310                 315                 320

Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile Ser Pro Ile
                325                 330                 335

Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly Gln Phe Leu
            340                 345                 350
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```
Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met Glu Ala Tyr
        355                 360                 365

Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg Met Lys Asn
    370                 375                 380

Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp Ser Glu Met
385                 390                 395                 400

Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe Ile Gln Ile
                405                 410                 415

Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His Tyr Ile Ala
            420                 425                 430

Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu Ala Pro Asp
        435                 440                 445

Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile
    450                 455                 460

Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr
465                 470                 475                 480

Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro
                485                 490                 495

Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn
            500                 505                 510

Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val
        515                 520                 525

Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys
    530                 535                 540

Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val
545                 550                 555                 560

Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg
                565                 570                 575

Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu
            580                 585                 590

Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly
        595                 600                 605

Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe
    610                 615                 620
```

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Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu
625                 630                 635                 640

Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser
                645                 650                 655

Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu
            660                 665                 670

Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly
        675                 680                 685

Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys
    690                 695                 700

His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly
705                 710                 715                 720

Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly
                725                 730                 735

Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys
            740                 745                 750

Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr
        755                 760                 765

Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu Pro
    770                 775                 780

Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His Gln Arg Glu
785                 790                 795                 800

Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp
                805                 810                 815

Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp
            820                 825                 830

Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His
        835                 840                 845

Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser
    850                 855                 860

Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln
865                 870                 875                 880

Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln
                885                 890                 895
```

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Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro
            900                 905                 910

Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn
        915                 920                 925

Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu
    930                 935                 940

Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro
945                 950                 955                 960

Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro
                965                 970                 975

Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val
            980                 985                 990

Asp Leu Glu Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
        995                 1000                1005

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1010                1015                1020

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1025                1030                1035

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1040                1045                1050

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1055                1060                1065

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1070                1075                1080

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1085                1090                1095

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1100                1105                1110

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1115                1120                1125

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1130                1135                1140

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1145                1150                1155
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Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1160                1165                1170

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1175                1180                1185

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1190                1195                1200

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1205                1210                1215

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1220                1225                1230

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1235                1240                1245

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1250                1255                1260

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1265                1270                1275

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1280                1285                1290

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1295                1300                1305

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1310                1315                1320

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1325                1330                1335

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1340                1345                1350

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1355                1360                1365

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1370                1375                1380

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1385                1390                1395

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1400                1405                1410
```

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1415 1420 1425

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1430 1435 1440

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1445 1450 1455

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1460 1465 1470

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1475 1480 1485

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1490 1495 1500

Ser Pro Ala Thr Gly His His His His His His
1505 1510

<210> 600
<211> 4545
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 600
gccaccagag gcgcgccagg ttctcctgct ggctccccca cctcaacaga agaggggaca 60

agcgaaagcg ctacgcctga gagtggccct ggctctgagc cagccacctc cggctctgaa 120

acccctgcct cgagcgctac aagaagatac tacctgggtg cagtggaact gtcatgggac 180

tatatgcaaa gtgatctcgg tgagctgcct gtggacgcaa gatttcctcc tagagtgcca 240

aaatcttttc cattcaacac ctcagtcgtg tacaaaaaga ctctgtttgt agaattcacg 300

gatcaccttt tcaacatcgc taagccaagg ccaccctgga tgggtctgct aggtcctacc 360

atccaggctg aggtttatga tacagtggtc attacactta agaacatggc ttcccatcct 420

gtcagtcttc atgctgttgg tgtatcctac tggaaagctt ctgagggagc tgaatatgat 480

gatcagacca gtcaaaggga gaaagaagat gataaagtct tccctggtgg aagccataca 540

tatgtctggc aggtcctgaa agagaatggt ccaatggcct ctgacccact gtgccttacc 600

tactcatatc tttctcatgt ggacctggta aaagacttga attcaggcct cattggagcc 660

ctactagtat gtagagaagg gagtctggcc aaggaaaaga cacagacctt gcacaaattt 720

atactacttt ttgctgtatt tgatgaaggg aaaagttggc actcagaaac aaagaactcc 780

ttgatgcagg atagggatgc tgcatctgct cgggcctggc ctaaaatgca cacagtcaat 840

ggttatgtaa acaggtctct gccaggtctg attggatgcc acaggaaatc agtctattgg 900

catgtgattg gaatgggcac cactcctgaa gtgcactcaa tattcctcga aggtcacaca 960

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caacatgatg gcatggaagc ttatgtcaaa gtagacagct gtccagagga accccaacta 1140

cgaatgaaaa ataatgaaga agcggaagac tatgatgatg atcttactga ttctgaaatg 1200

gatgtggtca ggtttgatga tgacaactct ccttccttta tccaaattcg ctcagttgcc 1260

aagaagcatc ctaaaacttg ggtacattac attgctgctg aagaggagga ctgggactat 1320

gctcccttag tcctcgcccc cgatgacaga agttataaaa gtcaatattt gaacaatggc 1380

cctcagcgga ttggtaggaa gtacaaaaaa gtccgattta tggcatacac agatgaaacc 1440

tttaagactc gtgaagctat tcagcatgaa tcaggaatct tgggaccttt actttatggg 1500

gaagttggag acacactgtt gattatattt aagaatcaag caagcagacc atataacatc 1560

taccctcacg gaatcactga tgtccgtcct ttgtattcaa ggagattacc aaaaggtgta 1620

aaacatttga aggattttcc aattctgcca ggagaaatat tcaaatataa atggacagtg 1680

actgtagaag atgggccaac taaatcagat cctcggtgcc tgacccgcta ttactctagt 1740

ttcgttaata tggagagaga tctagcttca ggactcattg gccctctcct catctgctac 1800

aaagaatctg tagatcaaag aggaaaccag ataatgtcag acaagaggaa tgtcatcctg 1860

ttttctgtat ttgatgagaa ccgaagctgg tacctcacag agaatataca acgctttctc 1920

cccaatccag ctggagtgca gcttgaggat ccagagttcc aagcctccaa catcatgcac 1980

agcatcaatg gctatgtttt tgatagtttg cagttgtcag tttgtttgca tgaggtggca 2040

tactggtaca ttctaagcat tggagcacag actgacttcc tttctgtctt cttctctgga 2100

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gaaactgtct tcatgtcgat ggaaaaccca ggtctatgga ttctggggtg ccacaactca 2220

gactttcgga acagaggcat gaccgcctta ctgaaggttt ctagttgtga caagaacact 2280

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ggcctgattg gacccttct ggtctgccac actaacacac tgaaccctgc tcatgggaga 3060

caagtgacag tacaggaatt tgctctgttt ttcaccatct ttgatgagac caaaagctgg 3120

tacttcactg aaaatatgga aagaaactgc agggctccct gcaatatcca gatggaagat 3180

cccactttta aagagaatta tcgcttccat gcaatcaatg gctacataat ggatacacta 3240

cctggcttag taatggctca ggatcaaagg attcgatggt atctgctcag catgggcagc 3300

aatgaaaaca tccattctat tcatttcagt ggacatgtgt tcactgtacg aaaaaaagag 3360

gagtataaaa tggcactgta caatctctat ccaggtgttt ttgagacagt ggaaatgtta 3420

ccatccaaag ctggaatttg gcgggtggaa tgccttattg gcgagcatct acatgctggg 3480

atgagcacac tttttctggt gtacagcaat aagtgtcaga ctcccctggg aatggcttct 3540

ggacacatta gagattttca gattacagct tcaggacaat atggacagtg ggccccaaag 3600

ctggccagac ttcattattc cggatcaatc aatgcctgga gcaccaagga gccctttct 3660

tggatcaagg tggatctgtt ggcaccaatg attattcacg gcatcaagac ccagggtgcc 3720

cgtcagaagt tctccagcct ctacatctct cagtttatca tcatgtatag tcttgatggg 3780

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cgtttgcacc caactcatta tagcattcgc agcactcttc gcatggagtt gatgggctgt 3960

gatttaaata gttgcagcat gccattggga atggagagta aagcaatatc agatgcacag 4020

attactgctt catcctactt taccaatatg tttgccacct ggtctccttc aaaagctcga 4080

cttcacctcc aagggaggag taatgcctgg agacctcagg tgaataatcc aaaagagtgg 4140

ctgcaagtgg acttccagaa gacaatgaaa gtcacaggag taactactca gggagtaaaa 4200

tctctgctta ccagcatgta tgtgaaggag ttcctcatct ccagcagtca agatggccat 4260

cagtggactc tctttttca gaatggcaaa gtaaaggttt ttcagggaaa tcaagactcc 4320

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ccccagagtt gggtgcacca gattgccctg aggatggagg ttctgggctg cgaggcacag 4440

gacctctacg gcgccggatc acctggggcc gaaacggccg aacaaaaact catctcagaa 4500

gaggatctgt cacctgcaac cggtcatcat caccatcacc attga 4545

<210> 601
<211> 1511
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 601
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr

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Met Gln Gly Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
20 25 30

Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
35 40 45

Pro Ala Thr Ser Gly Ser Glu Thr Pro Ala Ser Ser Ser Asp Leu Gly
50 55 60

Glu Leu Pro Val Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe
65 70 75 80

Pro Phe Asn Thr Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe
85 90 95

Thr Asp His Leu Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly
100 105 110

Leu Leu Gly Pro Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile
115 120 125

Thr Leu Lys Asn Met Ala Ser His Pro Val Ser Leu His Ala Val Gly
130 135 140

Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr
145 150 155 160

Ser Gln Arg Glu Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His
165 170 175

Thr Tyr Val Trp Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp
180 185 190

Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys
195 200 205

Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly
210 215 220

Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu
225 230 235 240

Phe Ala Val Phe Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn
245 250 255

Ser Leu Met Gln Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys
260 265 270

Met His Thr Val Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile

275 280 285

Gly Cys His Arg Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr
290 295 300

Thr Pro Glu Val His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val
305 310 315 320

Arg Asn His Arg Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu
325 330 335

Thr Ala Gln Thr Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys
340 345 350

His Ile Ser Ser His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val
355 360 365

Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu
370 375 380

Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val
385 390 395 400

Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val
405 410 415

Ala Lys Lys His Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu
420 425 430

Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser
435 440 445

Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys
450 455 460

Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr
465 470 475 480

Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr
485 490 495

Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser
500 505 510

Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu
515 520 525

Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro
530 535 540

Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu

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545                 550                 555                 560

Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser
                565                 570                 575

Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro
            580                 585                 590

Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile
        595                 600                 605

Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn
    610                 615                 620

Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro
625                 630                 635                 640

Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met
                645                 650                 655

His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys
            660                 665                 670

Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr
        675                 680                 685

Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met
    690                 695                 700

Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val
705                 710                 715                 720

Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn
                725                 730                 735

Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser
            740                 745                 750

Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile
        755                 760                 765

Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe
    770                 775                 780

Ser Gln Asn Pro Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg
785                 790                 795                 800

Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
                805                 810                 815

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu
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820 825 830

Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile
835 840 845

Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His
850 855 860

Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys
865 870 875 880

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
885 890 895

Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
900 905 910

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser
915 920 925

Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln
930 935 940

Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
945 950 955 960

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp
965 970 975

Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu
980 985 990

Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val Cys His Thr
995 1000 1005

Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr Val Gln Glu
1010 1015 1020

Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys Ser Trp Tyr
1025 1030 1035

Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro Cys Asn Ile
1040 1045 1050

Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg Phe His Ala
1055 1060 1065

Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu Val Met Ala
1070 1075 1080

Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met Gly Ser Asn

```
    1085                1090                1095

Glu Asn  Ile His Ser Ile His  Phe Ser Gly His Val  Phe Thr Val
    1100                1105                1110

Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
    1115                1120                1125

Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    1130                1135                1140

Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    1145                1150                1155

Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    1160                1165                1170

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
    1175                1180                1185

Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
    1190                1195                1200

Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
    1205                1210                1215

Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
    1220                1225                1230

Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    1235                1240                1245

Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
    1250                1255                1260

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    1265                1270                1275

Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
    1280                1285                1290

Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1295                1300                1305

Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
    1310                1315                1320

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    1325                1330                1335

Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
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1340 1345 1350

Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg
1355 1360 1365

Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln
1370 1375 1380

Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser
1385 1390 1395

Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser
1400 1405 1410

Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val
1415 1420 1425

Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn
1430 1435 1440

Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro
1445 1450 1455

Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly
1460 1465 1470

Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu
1475 1480 1485

Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala
1490 1495 1500

Thr Gly His His His His His His
1505 1510

<210> 602
<211> 4536
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 602
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ccaggttctc ctgctggctc ccccacctca acagaagagg ggacaagcga aagcgctacg 120

cctgagagtg gccctggctc tgagccagcc acctccggct ctgaaacccc tgcctcgagc 180

agtgatctcg gtgagctgcc tgtggacgca agatttcctc ctagagtgcc aaaatctttt 240

ccattcaaca cctcagtcgt gtacaaaaag actctgtttg tagaattcac ggatcacctt 300

ttcaacatcg ctaagccaag gccaccctgg atgggtctgc taggtcctac catccaggct 360

gaggtttatg atacagtggt cattacactt aagaacatgg cttcccatcc tgtcagtctt 420
catgctgttg gtgtatccta ctggaaagct tctgagggag ctgaatatga tgatcagacc 480
agtcaaaggg agaaagaaga tgataaagtc ttccctggtg gaagccatac atatgtctgg 540
caggtcctga aagagaatgg tccaatggcc tctgacccac tgtgccttac ctactcatat 600
ctttctcatg tggacctggt aaaagacttg aattcaggcc tcattggagc cctactagta 660
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tttgctgtat ttgatgaagg gaaaagttgg cactcagaaa caaagaactc cttgatgcag 780
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gacacactgt tgattatatt taagaatcaa gcaagcagac catataacat ctaccctcac 1560
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aagaaggaag attttgacat ttatgatgag gatgaaaatc agagcccccg cagctttcaa 2520

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gtaactttca gaaatcaggc ctctcgtccc tattccttct attctagcct tatttcttat 2820

gaggaagatc agaggcaagg agcagaacct agaaaaaact ttgtcaagcc taatgaaacc 2880

aaaacttact tttggaaagt gcaacatcat atggcaccca ctaaagatga gtttgactgc 2940

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aaagagaatt atcgcttcca tgcaatcaat ggctacataa tggatacact acctggctta 3240

gtaatggctc aggatcaaag gattcgatgg tatctgctca gcatgggcag caatgaaaac 3300

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cagacttatc gaggaaattc cactggaacc ttaatggtct tctttggcaa tgtggattca 3840

tctgggataa aacacaatat ttttaaccct ccaattattg ctcgatacat ccgtttgcac 3900

ccaactcatt atagcattcg cagcactctt cgcatggagt tgatgggctg tgatttaaat 3960

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tcatcctact ttaccaatat gtttgccacc tggtctcctt caaaagctcg acttcacctc 4080

caagggagga gtaatgcctg gagacctcag gtgaataatc caaaagagtg gctgcaagtg 4140

gacttccaga agacaatgaa agtcacagga gtaactactc agggagtaaa atctctgctt 4200

accagcatgt atgtgaagga gttcctcatc tccagcagtc aagatggcca tcagtggact 4260

ctctttttc agaatggcaa agtaaaggtt tttcagggaa atcaagactc cttcacacct 4320

gtggtgaact ctctagaccc accgttactg actcgctacc ttcgaattca cccccagagt 4380

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tcacctgcaa ccggtcatca tcaccatcac cattga 4536

<210> 603
<211> 1511
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 603
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Gly Ala Pro Gly Ser Pro Ala Gly Ser Pro
20 25 30

Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
35 40 45

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Ala Ser Ser
50 55 60

Glu Leu Pro Val Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe
65 70 75 80

Pro Phe Asn Thr Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe
85 90 95

Thr Asp His Leu Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly
100 105 110

Leu Leu Gly Pro Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile
115 120 125

Thr Leu Lys Asn Met Ala Ser His Pro Val Ser Leu His Ala Val Gly
130 135 140

Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr
145 150 155 160

Ser Gln Arg Glu Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His
165 170 175

Thr Tyr Val Trp Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp
180 185 190

Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys
195 200 205

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Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly
    210                 215                 220

Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu
225                 230                 235                 240

Phe Ala Val Phe Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn
                245                 250                 255

Ser Leu Met Gln Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys
            260                 265                 270

Met His Thr Val Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile
        275                 280                 285

Gly Cys His Arg Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr
    290                 295                 300

Thr Pro Glu Val His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val
305                 310                 315                 320

Arg Asn His Arg Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu
                325                 330                 335

Thr Ala Gln Thr Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys
            340                 345                 350

His Ile Ser Ser His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val
        355                 360                 365

Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu
    370                 375                 380

Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val
385                 390                 395                 400

Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val
                405                 410                 415

Ala Lys Lys His Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu
            420                 425                 430

Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser
        435                 440                 445

Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys
    450                 455                 460

Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr
465                 470                 475                 480
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Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr
                485                 490                 495

Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser
            500                 505                 510

Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu
        515                 520                 525

Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro
    530                 535                 540

Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu
545                 550                 555                 560

Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser
                565                 570                 575

Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro
            580                 585                 590

Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile
        595                 600                 605

Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn
    610                 615                 620

Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro
625                 630                 635                 640

Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met
                645                 650                 655

His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys
            660                 665                 670

Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr
        675                 680                 685

Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met
    690                 695                 700

Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val
705                 710                 715                 720

Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn
                725                 730                 735

Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser
            740                 745                 750
```

```
Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile
        755                 760                 765

Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe
    770                 775                 780

Ser Gln Asn Pro Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg
785                 790                 795                 800

Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
                805                 810                 815

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu
            820                 825                 830

Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile
        835                 840                 845

Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His
    850                 855                 860

Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys
865                 870                 875                 880

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
                885                 890                 895

Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
            900                 905                 910

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser
        915                 920                 925

Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln
    930                 935                 940

Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
945                 950                 955                 960

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp
                965                 970                 975

Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu
            980                 985                 990

Lys Asp Val His Ser Gly Leu Ile  Gly Pro Leu Leu Val  Cys His Thr
        995                 1000                1005

Asn Thr  Leu Asn Pro Ala His  Gly Arg Gln Val Thr  Val Gln Glu
    1010                1015                1020
```

```
Phe Ala  Leu Phe Phe Thr Ile  Phe Asp Glu Thr Lys  Ser Trp Tyr
    1025                1030                1035

Phe Thr  Glu Asn Met Glu Arg  Asn Cys Arg Ala Pro  Cys Asn Ile
    1040                1045                1050

Gln Met  Glu Asp Pro Thr Phe  Lys Glu Asn Tyr Arg  Phe His Ala
    1055                1060                1065

Ile Asn  Gly Tyr Ile Met Asp  Thr Leu Pro Gly Leu  Val Met Ala
    1070                1075                1080

Gln Asp  Gln Arg Ile Arg Trp  Tyr Leu Leu Ser Met  Gly Ser Asn
    1085                1090                1095

Glu Asn  Ile His Ser Ile His  Phe Ser Gly His Val  Phe Thr Val
    1100                1105                1110

Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
    1115                1120                1125

Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    1130                1135                1140

Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    1145                1150                1155

Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    1160                1165                1170

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
    1175                1180                1185

Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
    1190                1195                1200

Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
    1205                1210                1215

Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
    1220                1225                1230

Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    1235                1240                1245

Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
    1250                1255                1260

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    1265                1270                1275
```

```
Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
    1280                1285                1290

Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1295                1300                1305

Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
    1310                1315                1320

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    1325                1330                1335

Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
    1340                1345                1350

Ser Lys  Ala Arg Leu His Leu  Gln Gly Arg Ser Asn  Ala Trp Arg
    1355                1360                1365

Pro Gln  Val Asn Asn Pro Lys  Glu Trp Leu Gln Val  Asp Phe Gln
    1370                1375                1380

Lys Thr  Met Lys Val Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser
    1385                1390                1395

Leu Leu  Thr Ser Met Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser
    1400                1405                1410

Gln Asp  Gly His Gln Trp Thr  Leu Phe Phe Gln Asn  Gly Lys Val
    1415                1420                1425

Lys Val  Phe Gln Gly Asn Gln  Asp Ser Phe Thr Pro  Val Val Asn
    1430                1435                1440

Ser Leu  Asp Pro Pro Leu Leu  Thr Arg Tyr Leu Arg  Ile His Pro
    1445                1450                1455

Gln Ser  Trp Val His Gln Ile  Ala Leu Arg Met Glu  Val Leu Gly
    1460                1465                1470

Cys Glu  Ala Gln Asp Leu Tyr  Gly Ala Gly Ser Pro  Gly Ala Glu
    1475                1480                1485

Thr Ala  Glu Gln Lys Leu Ile  Ser Glu Glu Asp Leu  Ser Pro Ala
    1490                1495                1500

Thr Gly  His His His His His  His
    1505                1510
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<210> 604
<211> 4536
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

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gaaagcgcta cgcctgagag tggccctggc tctgagccag ccacctccgg ctctgaaacc 180

cctgcctcga gcgagctgcc tgtggacgca agatttcctc ctagagtgcc aaaatctttt 240

ccattcaaca cctcagtcgt gtacaaaaag actctgtttg tagaattcac ggatcacctt 300

ttcaacatcg ctaagccaag gccaccctgg atgggtctgc taggtcctac catccaggct 360

gaggtttatg atacagtggt cattacactt aagaacatgg cttcccatcc tgtcagtctt 420

catgctgttg gtgtatccta ctggaaagct tctgagggag ctgaatatga tgatcagacc 480

agtcaaaggg agaaagaaga tgataaagtc ttccctggtg gaagccatac atatgtctgg 540

caggtcctga aagagaatgg tccaatggcc tctgacccac tgtgccttac ctactcatat 600

ctttctcatg tggacctggt aaaagacttg aattcaggcc tcattggagc cctactagta 660

tgtagagaag ggagtctggc caaggaaaag acacagacct tgcacaaatt tatactactt 720

tttgctgtat ttgatgaagg gaaaagttgg cactcagaaa caaagaactc cttgatgcag 780

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aacaggtctc tgccaggtct gattggatgc cacaggaaat cagtctattg gcatgtgatt 900

ggaatgggca ccactcctga agtgcactca atattcctcg aaggtcacac atttcttgtg 960

aggaaccatc gccaggctag cttggaaatc tcgccaataa ctttccttac tgctcaaaca 1020

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ggcatggaag cttatgtcaa agtagacagc tgtccagagg aaccccaact acgaatgaaa 1140

aataatgaag aagcggaaga ctatgatgat gatcttactg attctgaaat ggatgtggtc 1200

aggtttgatg atgacaactc tccttccttt atccaaattc gctcagttgc caagaagcat 1260

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attggtagga agtacaaaaa agtccgattt atggcataca cagatgaaac ctttaagact 1440

cgtgaagcta ttcagcatga atcaggaatc ttgggacctt tactttatgg ggaagttgga 1500

gacacactgt tgattatatt taagaatcaa gcaagcagac catataacat ctaccctcac 1560

ggaatcactg atgtccgtcc tttgtattca aggagattac caaaaggtgt aaaacatttg 1620

aaggattttc caattctgcc aggagaaata ttcaaatata aatggacagt gactgtagaa 1680

gatgggccaa ctaaatcaga tcctcggtgc ctgacccgct attactctag tttcgttaat 1740

atggagagag atctagcttc aggactcatt ggccctctcc tcatctgcta caaagaatct 1800

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tttgatgaga accgaagctg gtacctcaca gagaatatac aacgctttct ccccaatcca 1920

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<210> 605
<211> 1511
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 605

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Gly Ser Pro
20 25 30

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr
35 40 45

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
50 55 60

Pro Ala Ser Ser Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe
65 70 75 80

Pro Phe Asn Thr Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe
85 90 95

Thr Asp His Leu Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly
100 105 110

Leu Leu Gly Pro Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile
115 120 125

Thr Leu Lys Asn Met Ala Ser His Pro Val Ser Leu His Ala Val Gly
130 135 140

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Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr
145                 150                 155                 160

Ser Gln Arg Glu Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His
                165                 170                 175

Thr Tyr Val Trp Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp
            180                 185                 190

Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys
        195                 200                 205

Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly
    210                 215                 220

Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu
225                 230                 235                 240

Phe Ala Val Phe Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn
                245                 250                 255

Ser Leu Met Gln Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys
            260                 265                 270

Met His Thr Val Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile
        275                 280                 285

Gly Cys His Arg Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr
    290                 295                 300

Thr Pro Glu Val His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val
305                 310                 315                 320

Arg Asn His Arg Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu
                325                 330                 335

Thr Ala Gln Thr Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys
            340                 345                 350

His Ile Ser Ser His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val
        355                 360                 365

Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu
    370                 375                 380

Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val
385                 390                 395                 400

Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val
                405                 410                 415
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Ala Lys Lys His Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu
            420                 425                 430

Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser
        435                 440                 445

Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys
    450                 455                 460

Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr
465                 470                 475                 480

Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr
                485                 490                 495

Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser
            500                 505                 510

Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu
        515                 520                 525

Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro
    530                 535                 540

Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu
545                 550                 555                 560

Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser
                565                 570                 575

Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro
            580                 585                 590

Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile
        595                 600                 605

Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn
    610                 615                 620

Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro
625                 630                 635                 640

Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met
                645                 650                 655

His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys
            660                 665                 670

Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr
        675                 680                 685
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Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met
    690                 695                 700

Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val
705                 710                 715                 720

Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn
                725                 730                 735

Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser
            740                 745                 750

Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile
        755                 760                 765

Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe
    770                 775                 780

Ser Gln Asn Pro Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg
785                 790                 795                 800

Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
                805                 810                 815

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu
            820                 825                 830

Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile
        835                 840                 845

Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His
    850                 855                 860

Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys
865                 870                 875                 880

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
                885                 890                 895

Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
            900                 905                 910

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser
        915                 920                 925

Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln
    930                 935                 940

Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
945                 950                 955                 960
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Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp
                965                 970                 975

Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu
            980                 985                 990

Lys Asp Val His Ser Gly Leu Ile  Gly Pro Leu Leu Val  Cys His Thr
        995                 1000                1005

Asn Thr  Leu Asn Pro Ala His  Gly Arg Gln Val Thr  Val Gln Glu
    1010                1015                1020

Phe Ala  Leu Phe Phe Thr Ile  Phe Asp Glu Thr Lys  Ser Trp Tyr
    1025                1030                1035

Phe Thr  Glu Asn Met Glu Arg  Asn Cys Arg Ala Pro  Cys Asn Ile
    1040                1045                1050

Gln Met  Glu Asp Pro Thr Phe  Lys Glu Asn Tyr Arg  Phe His Ala
    1055                1060                1065

Ile Asn  Gly Tyr Ile Met Asp  Thr Leu Pro Gly Leu  Val Met Ala
    1070                1075                1080

Gln Asp  Gln Arg Ile Arg Trp  Tyr Leu Leu Ser Met  Gly Ser Asn
    1085                1090                1095

Glu Asn  Ile His Ser Ile His  Phe Ser Gly His Val  Phe Thr Val
    1100                1105                1110

Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
    1115                1120                1125

Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    1130                1135                1140

Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    1145                1150                1155

Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    1160                1165                1170

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
    1175                1180                1185

Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
    1190                1195                1200

Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
    1205                1210                1215
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Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
    1220                1225                1230

Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    1235                1240                1245

Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
    1250                1255                1260

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    1265                1270                1275

Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
    1280                1285                1290

Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1295                1300                1305

Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
    1310                1315                1320

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    1325                1330                1335

Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
    1340                1345                1350

Ser Lys  Ala Arg Leu His Leu  Gln Gly Arg Ser Asn  Ala Trp Arg
    1355                1360                1365

Pro Gln  Val Asn Asn Pro Lys  Glu Trp Leu Gln Val  Asp Phe Gln
    1370                1375                1380

Lys Thr  Met Lys Val Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser
    1385                1390                1395

Leu Leu  Thr Ser Met Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser
    1400                1405                1410

Gln Asp  Gly His Gln Trp Thr  Leu Phe Phe Gln Asn  Gly Lys Val
    1415                1420                1425

Lys Val  Phe Gln Gly Asn Gln  Asp Ser Phe Thr Pro  Val Val Asn
    1430                1435                1440

Ser Leu  Asp Pro Pro Leu Leu  Thr Arg Tyr Leu Arg  Ile His Pro
    1445                1450                1455

Gln Ser  Trp Val His Gln Ile  Ala Leu Arg Met Glu  Val Leu Gly
    1460                1465                1470
```

Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu
1475 1480 1485

Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala
1490 1495 1500

Thr Gly His His His His His His
1505 1510

<210> 606
<211> 4536
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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gaggggacaa gcgaaagcgc tacgcctgag agtggccctg gctctgagcc agccacctcc 180

ggctctgaaa cccctgcctc gagcgacgca agatttcctc ctagagtgcc aaaatctttt 240

ccattcaaca cctcagtcgt gtacaaaaag actctgtttg tagaattcac ggatcacctt 300

ttcaacatcg ctaagccaag gccaccctgg atgggtctgc taggtcctac catccaggct 360

gaggtttatg atacagtggt cattacactt aagaacatgg cttcccatcc tgtcagtctt 420

catgctgttg gtgtatccta ctggaaagct tctgagggag ctgaatatga tgatcagacc 480

agtcaaaggg agaaagaaga tgataaagtc ttccctggtg gaagccatac atatgtctgg 540

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tgtagagaag ggagtctggc caaggaaaag acacagacct tgcacaaatt tatactactt 720

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ccaactcatt atagcattcg cagcactctt cgcatggagt tgatgggctg tgatttaaat 3960

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ctctttttc agaatggcaa agtaaaggtt tttcagggaa atcaagactc cttcacacct 4320

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<210> 607
<211> 1511
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 607
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Gly Ala Pro Gly Ser Pro Ala Gly
35 40 45

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
50 55 60

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Ala
65 70 75 80

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Ser Ser Asn Thr Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe
                85                  90                  95

Thr Asp His Leu Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly
            100                 105                 110

Leu Leu Gly Pro Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile
        115                 120                 125

Thr Leu Lys Asn Met Ala Ser His Pro Val Ser Leu His Ala Val Gly
    130                 135                 140

Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr
145                 150                 155                 160

Ser Gln Arg Glu Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His
                165                 170                 175

Thr Tyr Val Trp Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp
            180                 185                 190

Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys
        195                 200                 205

Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly
    210                 215                 220

Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu
225                 230                 235                 240

Phe Ala Val Phe Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn
                245                 250                 255

Ser Leu Met Gln Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys
            260                 265                 270

Met His Thr Val Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile
        275                 280                 285

Gly Cys His Arg Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr
    290                 295                 300

Thr Pro Glu Val His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val
305                 310                 315                 320

Arg Asn His Arg Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu
                325                 330                 335

Thr Ala Gln Thr Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys
            340                 345                 350
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```
His Ile Ser Ser His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val
        355                 360                 365

Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu
    370                 375                 380

Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val
385                 390                 395                 400

Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val
                405                 410                 415

Ala Lys Lys His Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu
            420                 425                 430

Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser
        435                 440                 445

Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys
    450                 455                 460

Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr
465                 470                 475                 480

Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr
                485                 490                 495

Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser
            500                 505                 510

Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu
        515                 520                 525

Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro
    530                 535                 540

Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu
545                 550                 555                 560

Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser
                565                 570                 575

Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro
            580                 585                 590

Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile
        595                 600                 605

Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn
    610                 615                 620
```

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Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro
625                 630                 635                 640

Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met
                645                 650                 655

His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys
            660                 665                 670

Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr
        675                 680                 685

Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met
    690                 695                 700

Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val
705                 710                 715                 720

Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn
                725                 730                 735

Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser
            740                 745                 750

Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile
        755                 760                 765

Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe
    770                 775                 780

Ser Gln Asn Pro Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg
785                 790                 795                 800

Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
                805                 810                 815

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu
            820                 825                 830

Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile
        835                 840                 845

Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His
    850                 855                 860

Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys
865                 870                 875                 880

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
                885                 890                 895
```

```
Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
            900                 905                 910

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser
        915                 920                 925

Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln
    930                 935                 940

Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
945                 950                 955                 960

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp
                965                 970                 975

Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu
            980                 985                 990

Lys Asp Val His Ser Gly Leu Ile  Gly Pro Leu Leu Val  Cys His Thr
        995                 1000                1005

Asn Thr  Leu Asn Pro Ala His  Gly Arg Gln Val Thr  Val Gln Glu
    1010                1015                1020

Phe Ala  Leu Phe Phe Thr Ile  Phe Asp Glu Thr Lys  Ser Trp Tyr
    1025                1030                1035

Phe Thr  Glu Asn Met Glu Arg  Asn Cys Arg Ala Pro  Cys Asn Ile
    1040                1045                1050

Gln Met  Glu Asp Pro Thr Phe  Lys Glu Asn Tyr Arg  Phe His Ala
    1055                1060                1065

Ile Asn  Gly Tyr Ile Met Asp  Thr Leu Pro Gly Leu  Val Met Ala
    1070                1075                1080

Gln Asp  Gln Arg Ile Arg Trp  Tyr Leu Leu Ser Met  Gly Ser Asn
    1085                1090                1095

Glu Asn  Ile His Ser Ile His  Phe Ser Gly His Val  Phe Thr Val
    1100                1105                1110

Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
    1115                1120                1125

Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    1130                1135                1140

Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    1145                1150                1155
```

```
Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    1160                1165                1170

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
    1175                1180                1185

Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
    1190                1195                1200

Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
    1205                1210                1215

Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
    1220                1225                1230

Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    1235                1240                1245

Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
    1250                1255                1260

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    1265                1270                1275

Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
    1280                1285                1290

Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1295                1300                1305

Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
    1310                1315                1320

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    1325                1330                1335

Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
    1340                1345                1350

Ser Lys  Ala Arg Leu His Leu  Gln Gly Arg Ser Asn  Ala Trp Arg
    1355                1360                1365

Pro Gln  Val Asn Asn Pro Lys  Glu Trp Leu Gln Val  Asp Phe Gln
    1370                1375                1380

Lys Thr  Met Lys Val Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser
    1385                1390                1395

Leu Leu  Thr Ser Met Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser
    1400                1405                1410
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Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val
1415 1420 1425

Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn
1430 1435 1440

Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro
1445 1450 1455

Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly
1460 1465 1470

Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu
1475 1480 1485

Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala
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Thr Gly His His His His His His
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<211> 4536
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ggcgcgccag gttctcctgc tggctccccc acctcaacag aagaggggac aagcgaaagc 180

gctacgcctg agagtggccc tggctctgag ccagccacct ccggctctga aacccctgcc 240

tcgagcaaca cctcagtcgt gtacaaaaag actctgtttg tagaattcac ggatcacctt 300

ttcaacatcg ctaagccaag gccaccctgg atgggtctgc taggtcctac catccaggct 360

gaggtttatg atacagtggt cattacactt aagaacatgg cttcccatcc tgtcagtctt 420

catgctgttg gtgtatccta ctggaaagct tctgagggag ctgaatatga tgatcagacc 480

agtcaaaggg agaaagaaga tgataaagtc ttccctggtg gaagccatac atatgtctgg 540

caggtcctga aagagaatgg tccaatggcc tctgacccac tgtgccttac ctactcatat 600

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tgtagagaag ggagtctggc caaggaaaag acacagacct tgcacaaatt tatactactt 720

tttgctgtat ttgatgaagg gaaaagttgg cactcagaaa caaagaactc cttgatgcag 780

gatagggatg ctgcatctgc tcgggcctgg cctaaaatgc acacagtcaa tggttatgta 840

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tcatcctact ttaccaatat gtttgccacc tggtctcctt caaaagctcg acttcacctc 4080

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<211> 1511
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Gly Ala Pro Gly
50 55 60

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
65 70 75 80

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
85 90 95

Glu Thr Pro Ala Ser Ser Ile Ala Lys Pro Arg Pro Pro Trp Met Gly
100 105 110

Leu Leu Gly Pro Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile
115 120 125

Thr Leu Lys Asn Met Ala Ser His Pro Val Ser Leu His Ala Val Gly
130 135 140

Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr
145 150 155 160

Ser Gln Arg Glu Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His
165 170 175

Thr Tyr Val Trp Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp
180 185 190

Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys
195 200 205

Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly
210 215 220

Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu
225 230 235 240

Phe Ala Val Phe Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn
245 250 255

Ser Leu Met Gln Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys
260 265 270

Met His Thr Val Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile

275 280 285

Gly Cys His Arg Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr
290 295 300

Thr Pro Glu Val His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val
305 310 315 320

Arg Asn His Arg Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu
325 330 335

Thr Ala Gln Thr Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys
340 345 350

His Ile Ser Ser His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val
355 360 365

Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu
370 375 380

Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val
385 390 395 400

Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val
405 410 415

Ala Lys Lys His Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu
420 425 430

Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser
435 440 445

Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys
450 455 460

Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr
465 470 475 480

Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr
485 490 495

Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser
500 505 510

Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu
515 520 525

Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro
530 535 540

Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu

545 550 555 560

Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser
565 570 575

Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro
580 585 590

Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile
595 600 605

Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn
610 615 620

Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro
625 630 635 640

Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met
645 650 655

His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys
660 665 670

Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr
675 680 685

Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met
690 695 700

Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val
705 710 715 720

Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn
725 730 735

Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser
740 745 750

Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile
755 760 765

Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe
770 775 780

Ser Gln Asn Pro Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg
785 790 795 800

Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
805 810 815

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu

820 825 830

Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile
835 840 845

Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His
850 855 860

Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys
865 870 875 880

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
885 890 895

Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
900 905 910

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser
915 920 925

Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln
930 935 940

Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
945 950 955 960

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp
965 970 975

Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu
980 985 990

Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val Cys His Thr
995 1000 1005

Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr Val Gln Glu
1010 1015 1020

Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys Ser Trp Tyr
1025 1030 1035

Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro Cys Asn Ile
1040 1045 1050

Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg Phe His Ala
1055 1060 1065

Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu Val Met Ala
1070 1075 1080

Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met Gly Ser Asn

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Glu Asn  Ile His Ser Ile His  Phe Ser Gly His Val  Phe Thr Val
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Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
    1115                1120                1125

Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    1130                1135                1140

Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    1145                1150                1155

Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    1160                1165                1170

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
    1175                1180                1185

Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
    1190                1195                1200

Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
    1205                1210                1215

Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
    1220                1225                1230

Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    1235                1240                1245

Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
    1250                1255                1260

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    1265                1270                1275

Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
    1280                1285                1290

Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1295                1300                1305

Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
    1310                1315                1320

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    1325                1330                1335

Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
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Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg
1355 1360 1365

Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln
1370 1375 1380

Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser
1385 1390 1395

Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser
1400 1405 1410

Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val
1415 1420 1425

Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn
1430 1435 1440

Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro
1445 1450 1455

Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly
1460 1465 1470

Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu
1475 1480 1485

Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala
1490 1495 1500

Thr Gly His His His His His His
1505 1510

<210> 610
<211> 4536
<212> DNA
<213> Artificial Sequence

<220>
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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

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<211> 1511
<212> PRT
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<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gly Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
115 120 125

Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
130 135 140

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Ala Ser Ser Gln Thr
145 150 155 160

Ser Gln Arg Glu Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His
165 170 175

Thr Tyr Val Trp Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp
180 185 190

Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys
195 200 205

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Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly
    210                 215                 220

Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu
225                 230                 235                 240

Phe Ala Val Phe Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn
                245                 250                 255

Ser Leu Met Gln Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys
            260                 265                 270

Met His Thr Val Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile
        275                 280                 285

Gly Cys His Arg Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr
    290                 295                 300

Thr Pro Glu Val His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val
305                 310                 315                 320

Arg Asn His Arg Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu
                325                 330                 335

Thr Ala Gln Thr Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys
            340                 345                 350

His Ile Ser Ser His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val
        355                 360                 365

Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu
    370                 375                 380

Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val
385                 390                 395                 400

Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val
                405                 410                 415

Ala Lys Lys His Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu
            420                 425                 430

Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser
        435                 440                 445

Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys
    450                 455                 460

Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr
465                 470                 475                 480
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Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr
                485                 490                 495

Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser
            500                 505                 510

Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu
        515                 520                 525

Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro
    530                 535                 540

Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu
545                 550                 555                 560

Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser
                565                 570                 575

Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro
            580                 585                 590

Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile
        595                 600                 605

Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn
    610                 615                 620

Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro
625                 630                 635                 640

Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met
                645                 650                 655

His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys
            660                 665                 670

Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr
        675                 680                 685

Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met
    690                 695                 700

Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val
705                 710                 715                 720

Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn
                725                 730                 735

Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser
            740                 745                 750
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```
Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile
        755                 760                 765

Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe
    770                 775                 780

Ser Gln Asn Pro Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg
785                 790                 795                 800

Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
                805                 810                 815

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu
            820                 825                 830

Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile
        835                 840                 845

Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His
    850                 855                 860

Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys
865                 870                 875                 880

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
                885                 890                 895

Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
            900                 905                 910

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser
        915                 920                 925

Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln
    930                 935                 940

Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
945                 950                 955                 960

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp
                965                 970                 975

Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu
            980                 985                 990

Lys Asp Val His Ser Gly Leu Ile  Gly Pro Leu Leu Val  Cys His Thr
        995                 1000                1005

Asn Thr  Leu Asn Pro Ala His  Gly Arg Gln Val Thr  Val Gln Glu
    1010                1015                1020
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Phe Ala  Leu Phe Phe Thr Ile  Phe Asp Glu Thr Lys  Ser Trp Tyr
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Phe Thr  Glu Asn Met Glu Arg  Asn Cys Arg Ala Pro  Cys Asn Ile
    1040                1045                1050

Gln Met  Glu Asp Pro Thr Phe  Lys Glu Asn Tyr Arg  Phe His Ala
    1055                1060                1065

Ile Asn  Gly Tyr Ile Met Asp  Thr Leu Pro Gly Leu  Val Met Ala
    1070                1075                1080

Gln Asp  Gln Arg Ile Arg Trp  Tyr Leu Leu Ser Met  Gly Ser Asn
    1085                1090                1095

Glu Asn  Ile His Ser Ile His  Phe Ser Gly His Val  Phe Thr Val
    1100                1105                1110

Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
    1115                1120                1125

Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    1130                1135                1140

Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    1145                1150                1155

Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    1160                1165                1170

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
    1175                1180                1185

Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
    1190                1195                1200

Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
    1205                1210                1215

Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
    1220                1225                1230

Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    1235                1240                1245

Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
    1250                1255                1260

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    1265                1270                1275
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Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
    1280                1285                1290

Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1295                1300                1305

Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
    1310                1315                1320

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    1325                1330                1335

Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
    1340                1345                1350

Ser Lys  Ala Arg Leu His Leu  Gln Gly Arg Ser Asn  Ala Trp Arg
    1355                1360                1365

Pro Gln  Val Asn Asn Pro Lys  Glu Trp Leu Gln Val  Asp Phe Gln
    1370                1375                1380

Lys Thr  Met Lys Val Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser
    1385                1390                1395

Leu Leu  Thr Ser Met Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser
    1400                1405                1410

Gln Asp  Gly His Gln Trp Thr  Leu Phe Phe Gln Asn  Gly Lys Val
    1415                1420                1425

Lys Val  Phe Gln Gly Asn Gln  Asp Ser Phe Thr Pro  Val Val Asn
    1430                1435                1440

Ser Leu  Asp Pro Pro Leu Leu  Thr Arg Tyr Leu Arg  Ile His Pro
    1445                1450                1455

Gln Ser  Trp Val His Gln Ile  Ala Leu Arg Met Glu  Val Leu Gly
    1460                1465                1470

Cys Glu  Ala Gln Asp Leu Tyr  Gly Ala Gly Ser Pro  Gly Ala Glu
    1475                1480                1485

Thr Ala  Glu Gln Lys Leu Ile  Ser Glu Glu Asp Leu  Ser Pro Ala
    1490                1495                1500

Thr Gly  His His His His His  His
    1505                1510
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<223> Description of Artificial Sequence: Synthetic
polynucleotide

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gtaactttca gaaatcaggc ctctcgtccc tattccttct attctagcct tatttcttat 2820

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gtaatggctc aggatcaaag gattcgatgg tatctgctca gcatgggcag caatgaaaac 3300

atccattcta ttcatttcag tggacatgtg ttcactgtac gaaaaaaaga ggagtataaa 3360

atggcactgt acaatctcta tccaggtgtt tttgagacag tggaaatgtt accatccaaa 3420

gctggaattt ggcgggtgga atgccttatt ggcgagcatc tacatgctgg gatgagcaca 3480

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gtggatctgt tggcaccaat gattattcac ggcatcaaga cccagggtgc ccgtcagaag 3720

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cagacttatc gaggaaattc cactggaacc ttaatggtct tctttggcaa tgtggattca 3840

tctgggataa aacacaatat ttttaaccct ccaattattg ctcgatacat ccgtttgcac 3900

ccaactcatt atagcattcg cagcactctt cgcatggagt tgatgggctg tgatttaaat 3960

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tcatcctact ttaccaatat gtttgccacc tggtctcctt caaaagctcg acttcacctc 4080

caagggagga gtaatgcctg gagacctcag gtgaataatc caaaagagtg gctgcaagtg 4140

gacttccaga agacaatgaa agtcacagga gtaactactc agggagtaaa atctctgctt 4200

accagcatgt atgtgaagga gttcctcatc tccagcagtc aagatggcca tcagtggact 4260

ctctttttc agaatggcaa agtaaaggtt tttcagggaa atcaagactc cttcacacct 4320

gtggtgaact ctctagaccc accgttactg actcgctacc ttcgaattca cccccagagt 4380

tgggtgcacc agattgccct gaggatggag gttctgggct gcgaggcaca ggacctctac 4440

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tcacctgcaa ccggtcatca tcaccatcac cattga 4536

<210> 613
<211> 1511
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 613

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
130 135 140

```
Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
145                 150                 155                 160

Pro Ala Thr Ser Gly Ser Glu Thr Pro Ala Ser Ser Gly Gly Ser His
                165                 170                 175

Thr Tyr Val Trp Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp
            180                 185                 190

Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys
        195                 200                 205

Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly
    210                 215                 220

Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu
225                 230                 235                 240

Phe Ala Val Phe Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn
                245                 250                 255

Ser Leu Met Gln Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys
            260                 265                 270

Met His Thr Val Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile
        275                 280                 285

Gly Cys His Arg Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr
    290                 295                 300

Thr Pro Glu Val His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val
305                 310                 315                 320

Arg Asn His Arg Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu
                325                 330                 335

Thr Ala Gln Thr Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys
            340                 345                 350

His Ile Ser Ser His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val
        355                 360                 365

Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu
    370                 375                 380

Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val
385                 390                 395                 400

Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val
                405                 410                 415
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```
Ala Lys Lys His Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu
            420                 425                 430

Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser
        435                 440                 445

Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys
    450                 455                 460

Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr
465                 470                 475                 480

Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr
                485                 490                 495

Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser
            500                 505                 510

Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu
        515                 520                 525

Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro
    530                 535                 540

Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu
545                 550                 555                 560

Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser
                565                 570                 575

Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro
            580                 585                 590

Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile
        595                 600                 605

Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn
    610                 615                 620

Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro
625                 630                 635                 640

Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met
                645                 650                 655

His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys
            660                 665                 670

Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr
        675                 680                 685
```

```
Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met
    690                 695                 700

Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val
705                 710                 715                 720

Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn
                725                 730                 735

Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser
            740                 745                 750

Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile
        755                 760                 765

Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe
    770                 775                 780

Ser Gln Asn Pro Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg
785                 790                 795                 800

Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
                805                 810                 815

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu
            820                 825                 830

Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile
        835                 840                 845

Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His
    850                 855                 860

Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys
865                 870                 875                 880

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
                885                 890                 895

Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
            900                 905                 910

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser
        915                 920                 925

Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln
    930                 935                 940

Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
945                 950                 955                 960
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Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp
                965                 970                 975

Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu
            980                 985                 990

Lys Asp Val His Ser Gly Leu Ile  Gly Pro Leu Leu Val  Cys His Thr
        995                 1000                1005

Asn Thr  Leu Asn Pro Ala His  Gly Arg Gln Val Thr  Val Gln Glu
    1010                1015                1020

Phe Ala  Leu Phe Phe Thr Ile  Phe Asp Glu Thr Lys  Ser Trp Tyr
    1025                1030                1035

Phe Thr  Glu Asn Met Glu Arg  Asn Cys Arg Ala Pro  Cys Asn Ile
    1040                1045                1050

Gln Met  Glu Asp Pro Thr Phe  Lys Glu Asn Tyr Arg  Phe His Ala
    1055                1060                1065

Ile Asn  Gly Tyr Ile Met Asp  Thr Leu Pro Gly Leu  Val Met Ala
    1070                1075                1080

Gln Asp  Gln Arg Ile Arg Trp  Tyr Leu Leu Ser Met  Gly Ser Asn
    1085                1090                1095

Glu Asn  Ile His Ser Ile His  Phe Ser Gly His Val  Phe Thr Val
    1100                1105                1110

Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
    1115                1120                1125

Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    1130                1135                1140

Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    1145                1150                1155

Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    1160                1165                1170

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
    1175                1180                1185

Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
    1190                1195                1200

Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
    1205                1210                1215
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```
Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
    1220                1225                1230

Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    1235                1240                1245

Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
    1250                1255                1260

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    1265                1270                1275

Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
    1280                1285                1290

Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1295                1300                1305

Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
    1310                1315                1320

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    1325                1330                1335

Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
    1340                1345                1350

Ser Lys  Ala Arg Leu His Leu  Gln Gly Arg Ser Asn  Ala Trp Arg
    1355                1360                1365

Pro Gln  Val Asn Asn Pro Lys  Glu Trp Leu Gln Val  Asp Phe Gln
    1370                1375                1380

Lys Thr  Met Lys Val Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser
    1385                1390                1395

Leu Leu  Thr Ser Met Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser
    1400                1405                1410

Gln Asp  Gly His Gln Trp Thr  Leu Phe Phe Gln Asn  Gly Lys Val
    1415                1420                1425

Lys Val  Phe Gln Gly Asn Gln  Asp Ser Phe Thr Pro  Val Val Asn
    1430                1435                1440

Ser Leu  Asp Pro Pro Leu Leu  Thr Arg Tyr Leu Arg  Ile His Pro
    1445                1450                1455

Gln Ser  Trp Val His Gln Ile  Ala Leu Arg Met Glu  Val Leu Gly
    1460                1465                1470
```

Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu
1475 1480 1485

Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala
1490 1495 1500

Thr Gly His His His His His His
1505 1510

<210> 614
<211> 4536
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 614
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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggcgcgccag gttctcctgc tggctccccc 420

acctcaacag aagaggggac aagcgaaagc gctacgcctg agagtggccc tggctctgag 480

ccagccacct ccggctctga aacccctgcc tcgagcggtg gaagccatac atatgtctgg 540

caggtcctga aagagaatgg tccaatggcc tctgacccac tgtgccttac ctactcatat 600

ctttctcatg tggacctggt aaaagacttg aattcaggcc tcattggagc cctactagta 660

tgtagagaag ggagtctggc caaggaaaag acacagacct tgcacaaatt tatactactt 720

tttgctgtat ttgatgaagg gaaaagttgg cactcagaaa caaagaactc cttgatgcag 780

gatagggatg ctgcatctgc tcgggcctgg cctaaaatgc acacagtcaa tggttatgta 840

aacaggtctc tgccaggtct gattggatgc cacaggaaat cagtctattg gcatgtgatt 900

ggaatgggca ccactcctga agtgcactca atattcctcg aaggtcacac atttcttgtg 960

aggaaccatc gccaggctag cttggaaatc tcgccaataa ctttccttac tgctcaaaca 1020

ctcttgatgg accttggaca gtttctactg ttttgtcata tctcttccca ccaacatgat 1080

ggcatggaag cttatgtcaa agtagacagc tgtccagagg aaccccaact acgaatgaaa 1140

aataatgaag aagcggaaga ctatgatgat gatcttactg attctgaaat ggatgtggtc 1200

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ccaactcatt atagcattcg cagcactctt cgcatggagt tgatgggctg tgatttaaat 3960

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tcatcctact ttaccaatat gtttgccacc tggtctcctt caaaagctcg acttcacctc 4080

caagggagga gtaatgcctg gagacctcag gtgaataatc caaaagagtg gctgcaagtg 4140

gacttccaga agacaatgaa agtcacagga gtaactactc agggagtaaa atctctgctt 4200

accagcatgt atgtgaagga gttcctcatc tccagcagtc aagatggcca tcagtggact 4260

ctctttttc agaatggcaa agtaaaggtt tttcagggaa atcaagactc cttcacacct 4320

gtggtgaact ctctagaccc accgttactg actcgctacc ttcgaattca cccccagagt 4380

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<210> 615
<211> 1511
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 615
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

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Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Gly Ala Pro Gly
            180                 185                 190

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
        195                 200                 205

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
    210                 215                 220

Glu Thr Pro Ala Ser Ser Thr Gln Thr Leu His Lys Phe Ile Leu Leu
225                 230                 235                 240

Phe Ala Val Phe Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn
                245                 250                 255

Ser Leu Met Gln Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys
            260                 265                 270

Met His Thr Val Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile
        275                 280                 285

Gly Cys His Arg Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr
    290                 295                 300

Thr Pro Glu Val His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val
305                 310                 315                 320

Arg Asn His Arg Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu
                325                 330                 335

Thr Ala Gln Thr Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys
            340                 345                 350
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His Ile Ser Ser His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val
        355                 360                 365

Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu
    370                 375                 380

Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val
385                 390                 395                 400

Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val
                405                 410                 415

Ala Lys Lys His Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu
            420                 425                 430

Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser
        435                 440                 445

Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys
    450                 455                 460

Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr
465                 470                 475                 480

Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr
                485                 490                 495

Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser
            500                 505                 510

Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu
        515                 520                 525

Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro
    530                 535                 540

Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu
545                 550                 555                 560

Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser
                565                 570                 575

Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro
            580                 585                 590

Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile
        595                 600                 605

Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn
    610                 615                 620
```

```
Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro
625                 630                 635                 640

Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met
                645                 650                 655

His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys
            660                 665                 670

Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr
        675                 680                 685

Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met
    690                 695                 700

Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val
705                 710                 715                 720

Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn
                725                 730                 735

Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser
            740                 745                 750

Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile
        755                 760                 765

Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe
    770                 775                 780

Ser Gln Asn Pro Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg
785                 790                 795                 800

Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
                805                 810                 815

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu
            820                 825                 830

Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile
        835                 840                 845

Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His
    850                 855                 860

Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys
865                 870                 875                 880

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
                885                 890                 895
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Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
            900                 905                 910

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser
        915                 920                 925

Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln
    930                 935                 940

Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
945                 950                 955                 960

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp
                965                 970                 975

Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu
            980                 985                 990

Lys Asp Val His Ser Gly Leu Ile  Gly Pro Leu Leu Val  Cys His Thr
        995                 1000                1005

Asn Thr  Leu Asn Pro Ala His  Gly Arg Gln Val Thr  Val Gln Glu
    1010                1015                1020

Phe Ala  Leu Phe Phe Thr Ile  Phe Asp Glu Thr Lys  Ser Trp Tyr
    1025                1030                1035

Phe Thr  Glu Asn Met Glu Arg  Asn Cys Arg Ala Pro  Cys Asn Ile
    1040                1045                1050

Gln Met  Glu Asp Pro Thr Phe  Lys Glu Asn Tyr Arg  Phe His Ala
    1055                1060                1065

Ile Asn  Gly Tyr Ile Met Asp  Thr Leu Pro Gly Leu  Val Met Ala
    1070                1075                1080

Gln Asp  Gln Arg Ile Arg Trp  Tyr Leu Leu Ser Met  Gly Ser Asn
    1085                1090                1095

Glu Asn  Ile His Ser Ile His  Phe Ser Gly His Val  Phe Thr Val
    1100                1105                1110

Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
    1115                1120                1125

Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    1130                1135                1140

Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    1145                1150                1155
```

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Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    1160                 1165                 1170

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
    1175                 1180                 1185

Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
    1190                 1195                 1200

Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
    1205                 1210                 1215

Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
    1220                 1225                 1230

Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    1235                 1240                 1245

Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
    1250                 1255                 1260

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    1265                 1270                 1275

Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
    1280                 1285                 1290

Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1295                 1300                 1305

Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
    1310                 1315                 1320

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    1325                 1330                 1335

Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
    1340                 1345                 1350

Ser Lys  Ala Arg Leu His Leu  Gln Gly Arg Ser Asn  Ala Trp Arg
    1355                 1360                 1365

Pro Gln  Val Asn Asn Pro Lys  Glu Trp Leu Gln Val  Asp Phe Gln
    1370                 1375                 1380

Lys Thr  Met Lys Val Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser
    1385                 1390                 1395

Leu Leu  Thr Ser Met Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser
    1400                 1405                 1410
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Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val
1415 1420 1425

Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn
1430 1435 1440

Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro
1445 1450 1455

Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly
1460 1465 1470

Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu
1475 1480 1485

Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala
1490 1495 1500

Thr Gly His His His His His His
1505 1510

<210> 616
<211> 4536
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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ctcggtgagc tgcctgtgga cgcaagattt cctcctagag tgccaaaatc ttttccattc 120

aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagggcgcg ccaggttctc ctgctggctc ccccacctca 600

acagaagagg ggacaagcga aagcgctacg cctgagagtg gccctggctc tgagccagcc 660

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tttgctgtat ttgatgaagg gaaaagttgg cactcagaaa caaagaactc cttgatgcag 780

gatagggatg ctgcatctgc tcgggcctgg cctaaaatgc acacagtcaa tggttatgta 840

aacaggtctc tgccaggtct gattggatgc cacaggaaat cagtctattg gcatgtgatt 900

ggaatgggca ccactcctga agtgcactca atattcctcg aaggtcacac atttcttgtg 960

aggaaccatc gccaggctag cttggaaatc tcgccaataa ctttccttac tgctcaaaca 1020

ctcttgatgg accttggaca gtttctactg ttttgtcata tctcttccca ccaacatgat 1080

ggcatggaag cttatgtcaa agtagacagc tgtccagagg aaccccaact acgaatgaaa 1140

aataatgaag aagcggaaga ctatgatgat gatcttactg attctgaaat ggatgtggtc 1200

aggtttgatg atgacaactc tccttccttt atccaaattc gctcagttgc caagaagcat 1260

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gtcctcgccc ccgatgacag aagttataaa agtcaatatt tgaacaatgg ccctcagcgg 1380

attggtagga agtacaaaaa agtccgattt atggcataca cagatgaaac ctttaagact 1440

cgtgaagcta ttcagcatga atcaggaatc ttgggacctt tactttatgg ggaagttgga 1500

gacacactgt tgattatatt taagaatcaa gcaagcagac catataacat ctaccctcac 1560

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gtagatcaaa gaggaaacca gataatgtca gacaagagga atgtcatcct gttttctgta 1860

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aacagaggca tgaccgcctt actgaaggtt tctagttgtg acaagaacac tggtgattat 2280

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actactcttc agtcagatca agaggaaatc gattatgatg ataccatatc agttgaaatg 2460

aagaaggaag attttgacat ttatgatgag gatgaaaatc agagcccccg cagctttcaa 2520

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ccaactcatt atagcattcg cagcactctt cgcatggagt tgatgggctg tgatttaaat 3960

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tcatcctact ttaccaatat gtttgccacc tggtctcctt caaaagctcg acttcacctc 4080

caagggagga gtaatgcctg gagacctcag gtgaataatc caaaagagtg gctgcaagtg 4140

gacttccaga agacaatgaa agtcacagga gtaactactc agggagtaaa atctctgctt 4200

accagcatgt atgtgaagga gttcctcatc tccagcagtc aagatggcca tcagtggact 4260

ctctttttc agaatggcaa agtaaaggtt tttcagggaa atcaagactc cttcacacct 4320

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tcacctgcaa ccggtcatca tcaccatcac cattga 4536

<210> 617
<211> 1511
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 617
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Gly Ala Pro Gly Ser Pro Ala Gly
210 215 220

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
225 230 235 240

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Ala
245 250 255

Ser Ser Met Gln Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys
260 265 270

Met His Thr Val Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile

275 280 285

Gly Cys His Arg Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr
290 295 300

Thr Pro Glu Val His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val
305 310 315 320

Arg Asn His Arg Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu
325 330 335

Thr Ala Gln Thr Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys
340 345 350

His Ile Ser Ser His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val
355 360 365

Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu
370 375 380

Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val
385 390 395 400

Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val
405 410 415

Ala Lys Lys His Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu
420 425 430

Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser
435 440 445

Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys
450 455 460

Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr
465 470 475 480

Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr
485 490 495

Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser
500 505 510

Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu
515 520 525

Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro
530 535 540

Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu

545 550 555 560

Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser
565 570 575

Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro
580 585 590

Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile
595 600 605

Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn
610 615 620

Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro
625 630 635 640

Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met
645 650 655

His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys
660 665 670

Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr
675 680 685

Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met
690 695 700

Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val
705 710 715 720

Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn
725 730 735

Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser
740 745 750

Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile
755 760 765

Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe
770 775 780

Ser Gln Asn Pro Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg
785 790 795 800

Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
805 810 815

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu

820 825 830

Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile
835 840 845

Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His
850 855 860

Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys
865 870 875 880

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
885 890 895

Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
900 905 910

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser
915 920 925

Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln
930 935 940

Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
945 950 955 960

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp
965 970 975

Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu
980 985 990

Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val Cys His Thr
995 1000 1005

Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr Val Gln Glu
1010 1015 1020

Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys Ser Trp Tyr
1025 1030 1035

Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro Cys Asn Ile
1040 1045 1050

Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg Phe His Ala
1055 1060 1065

Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu Val Met Ala
1070 1075 1080

Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met Gly Ser Asn

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Glu Asn  Ile His Ser Ile His  Phe Ser Gly His Val  Phe Thr Val
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Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
    1115                1120                1125

Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    1130                1135                1140

Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    1145                1150                1155

Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    1160                1165                1170

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
    1175                1180                1185

Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
    1190                1195                1200

Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
    1205                1210                1215

Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
    1220                1225                1230

Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    1235                1240                1245

Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
    1250                1255                1260

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    1265                1270                1275

Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
    1280                1285                1290

Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1295                1300                1305

Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
    1310                1315                1320

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    1325                1330                1335

Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
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1340 1345 1350

Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg
1355 1360 1365

Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln
1370 1375 1380

Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser
1385 1390 1395

Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser
1400 1405 1410

Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val
1415 1420 1425

Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn
1430 1435 1440

Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro
1445 1450 1455

Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly
1460 1465 1470

Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu
1475 1480 1485

Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala
1490 1495 1500

Thr Gly His His His His His His
1505 1510

<210> 618
<211> 4536
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

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gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

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<210> 619
<211> 1511
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 619

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

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His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Gly Ala Pro Gly Ser Pro Ala Gly Ser Pro
225                 230                 235                 240

Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
                245                 250                 255

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Ala Ser Ser
            260                 265                 270

Met His Thr Val Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile
        275                 280                 285

Gly Cys His Arg Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr
    290                 295                 300

Thr Pro Glu Val His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val
305                 310                 315                 320

Arg Asn His Arg Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu
                325                 330                 335

Thr Ala Gln Thr Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys
            340                 345                 350

His Ile Ser Ser His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val
        355                 360                 365

Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu
    370                 375                 380

Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val
385                 390                 395                 400

Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val
                405                 410                 415

Ala Lys Lys His Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu
            420                 425                 430

Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser
        435                 440                 445

Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys
    450                 455                 460

Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr
465                 470                 475                 480
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Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr
                485                 490                 495

Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser
            500                 505                 510

Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu
        515                 520                 525

Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro
    530                 535                 540

Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu
545                 550                 555                 560

Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser
                565                 570                 575

Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro
            580                 585                 590

Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile
        595                 600                 605

Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn
    610                 615                 620

Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro
625                 630                 635                 640

Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met
                645                 650                 655

His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys
            660                 665                 670

Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr
        675                 680                 685

Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met
    690                 695                 700

Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val
705                 710                 715                 720

Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn
                725                 730                 735

Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser
            740                 745                 750
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Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile
        755                 760                 765

Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe
    770                 775                 780

Ser Gln Asn Pro Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg
785                 790                 795                 800

Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
                805                 810                 815

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu
            820                 825                 830

Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile
        835                 840                 845

Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His
    850                 855                 860

Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys
865                 870                 875                 880

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
                885                 890                 895

Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
            900                 905                 910

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser
        915                 920                 925

Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln
    930                 935                 940

Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
945                 950                 955                 960

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp
                965                 970                 975

Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu
            980                 985                 990

Lys Asp Val His Ser Gly Leu Ile  Gly Pro Leu Leu Val  Cys His Thr
        995                 1000                1005

Asn Thr  Leu Asn Pro Ala His  Gly Arg Gln Val Thr  Val Gln Glu
    1010                1015                1020
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Phe Ala  Leu Phe Phe Thr Ile  Phe Asp Glu Thr Lys  Ser Trp Tyr
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Phe Thr  Glu Asn Met Glu Arg  Asn Cys Arg Ala Pro  Cys Asn Ile
    1040                1045                1050

Gln Met  Glu Asp Pro Thr Phe  Lys Glu Asn Tyr Arg  Phe His Ala
    1055                1060                1065

Ile Asn  Gly Tyr Ile Met Asp  Thr Leu Pro Gly Leu  Val Met Ala
    1070                1075                1080

Gln Asp  Gln Arg Ile Arg Trp  Tyr Leu Leu Ser Met  Gly Ser Asn
    1085                1090                1095

Glu Asn  Ile His Ser Ile His  Phe Ser Gly His Val  Phe Thr Val
    1100                1105                1110

Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
    1115                1120                1125

Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    1130                1135                1140

Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    1145                1150                1155

Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    1160                1165                1170

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
    1175                1180                1185

Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
    1190                1195                1200

Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
    1205                1210                1215

Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
    1220                1225                1230

Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    1235                1240                1245

Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
    1250                1255                1260

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    1265                1270                1275
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Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
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Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1295                1300                1305

Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
    1310                1315                1320

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    1325                1330                1335

Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
    1340                1345                1350

Ser Lys  Ala Arg Leu His Leu  Gln Gly Arg Ser Asn  Ala Trp Arg
    1355                1360                1365

Pro Gln  Val Asn Asn Pro Lys  Glu Trp Leu Gln Val  Asp Phe Gln
    1370                1375                1380

Lys Thr  Met Lys Val Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser
    1385                1390                1395

Leu Leu  Thr Ser Met Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser
    1400                1405                1410

Gln Asp  Gly His Gln Trp Thr  Leu Phe Phe Gln Asn  Gly Lys Val
    1415                1420                1425

Lys Val  Phe Gln Gly Asn Gln  Asp Ser Phe Thr Pro  Val Val Asn
    1430                1435                1440

Ser Leu  Asp Pro Pro Leu Leu  Thr Arg Tyr Leu Arg  Ile His Pro
    1445                1450                1455

Gln Ser  Trp Val His Gln Ile  Ala Leu Arg Met Glu  Val Leu Gly
    1460                1465                1470

Cys Glu  Ala Gln Asp Leu Tyr  Gly Ala Gly Ser Pro  Gly Ala Glu
    1475                1480                1485

Thr Ala  Glu Gln Lys Leu Ile  Ser Glu Glu Asp Leu  Ser Pro Ala
    1490                1495                1500

Thr Gly  His His His His His  His
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<210> 620
<211> 4536
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

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<210> 621
<211> 1514
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 621

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

```
Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gly Ala Pro
                325                 330                 335

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
            340                 345                 350

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
        355                 360                 365

Ser Glu Thr Pro Ala Ser Ser Val Gln Leu Gln Leu Arg Met Lys Asn
    370                 375                 380

Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp Ser Glu Met
385                 390                 395                 400

Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe Ile Gln Ile
                405                 410                 415
```

Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His Tyr Ile Ala
420 425 430

Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu Ala Pro Asp
435 440 445

Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile
450 455 460

Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr
465 470 475 480

Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro
485 490 495

Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn
500 505 510

Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val
515 520 525

Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys
530 535 540

Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val
545 550 555 560

Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg
565 570 575

Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu
580 585 590

Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly
595 600 605

Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe
610 615 620

Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu
625 630 635 640

Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser
645 650 655

Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu
660 665 670

Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly
675 680 685

```
Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys
    690                 695                 700

His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly
705                 710                 715                 720

Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly
                725                 730                 735

Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys
            740                 745                 750

Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr
        755                 760                 765

Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu Pro
    770                 775                 780

Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His Gln Arg Glu
785                 790                 795                 800

Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp
                805                 810                 815

Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp
            820                 825                 830

Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His
        835                 840                 845

Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser
    850                 855                 860

Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln
865                 870                 875                 880

Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln
                885                 890                 895

Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro
            900                 905                 910

Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn
        915                 920                 925

Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu
    930                 935                 940

Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro
945                 950                 955                 960
```

```
Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro
                965                 970                 975

Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val
            980                 985                 990

Asp Leu Glu Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
        995                 1000                1005

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1010                1015                1020

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1025                1030                1035

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1040                1045                1050

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1055                1060                1065

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1070                1075                1080

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1085                1090                1095

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1100                1105                1110

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1115                1120                1125

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1130                1135                1140

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1145                1150                1155

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1160                1165                1170

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1175                1180                1185

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1190                1195                1200

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1205                1210                1215
```

```
Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1220                1225                1230

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1235                1240                1245

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1250                1255                1260

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1265                1270                1275

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1280                1285                1290

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1295                1300                1305

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1310                1315                1320

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1325                1330                1335

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1340                1345                1350

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1355                1360                1365

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1370                1375                1380

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1385                1390                1395

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1400                1405                1410

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1415                1420                1425

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1430                1435                1440

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1445                1450                1455

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1460                1465                1470
```

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1475 1480 1485

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1490 1495 1500

Ser Pro Ala Thr Gly His His His His His His
1505 1510

<210> 622
<211> 4545
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 622
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ctcggtgagc tgcctgtgga cgcaagattt cctcctagag tgccaaaatc ttttccattc 120

aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccg gcgcgccagg ttctcctgct 1020

ggctccccca cctcaacaga agaggggaca agcgaaagcg ctacgcctga gagtggccct 1080

ggctctgagc cagccacctc cggctctgaa accctgcct cgagcgtgca acttcaacta 1140

cgaatgaaaa ataatgaaga agcggaagac tatgatgatg atcttactga ttctgaaatg 1200

gatgtggtca ggtttgatga tgacaactct ccttccttta tccaaattcg ctcagttgcc 1260

aagaagcatc ctaaaacttg ggtacattac attgctgctg aagaggagga ctgggactat 1320

gctcccttag tcctcgcccc cgatgacaga agttataaaa gtcaatattt gaacaatggc 1380

cctcagcgga ttggtaggaa gtacaaaaaa gtccgattta tggcatacac agatgaaacc 1440

tttaagactc gtgaagctat tcagcatgaa tcaggaatct tgggaccttt actttatggg 1500

gaagttggag acacactgtt gattatattt aagaatcaag caagcagacc atataacatc 1560

taccctcacg gaatcactga tgtccgtcct ttgtattcaa ggagattacc aaaaggtgta 1620

aaacatttga aggattttcc aattctgcca ggagaaatat tcaaatataa atggacagtg 1680

actgtagaag atgggccaac taaatcagat cctcggtgcc tgacccgcta ttactctagt 1740

ttcgttaata tggagagaga tctagcttca ggactcattg gccctctcct catctgctac 1800

aaagaatctg tagatcaaag aggaaaccag ataatgtcag acaagaggaa tgtcatcctg 1860

ttttctgtat ttgatgagaa ccgaagctgg tacctcacag agaatataca acgctttctc 1920

cccaatccag ctggagtgca gcttgaggat ccagagttcc aagcctccaa catcatgcac 1980

agcatcaatg gctatgtttt tgatagtttg cagttgtcag tttgtttgca tgaggtggca 2040

tactggtaca ttctaagcat tggagcacag actgacttcc tttctgtctt cttctctgga 2100

tataccttca aacacaaaat ggtctatgaa gacacactca ccctattccc attctcagga 2160

gaaactgtct tcatgtcgat ggaaaaccca ggtctatgga ttctggggtg ccacaactca 2220

gactttcgga acagaggcat gaccgcctta ctgaaggttt ctagttgtga caagaacact 2280

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gggatgagta gctccccaca tgttctaaga aacagggctc agagtggcag tgtccctcag 2640

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ggagaactaa atgaacattt gggactcctg gggccatata taagagcaga agttgaagat 2760

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ggcctgattg gaccccttct ggtctgccac actaacacac tgaaccctgc tcatgggaga 3060

caagtgacag tacaggaatt tgctctgttt ttcaccatct ttgatgagac caaaagctgg 3120

tacttcactg aaaatatgga aagaaactgc agggctccct gcaatatcca gatggaagat 3180

cccactttta aagagaatta tcgcttccat gcaatcaatg gctacataat ggatacacta 3240

cctggcttag taatggctca ggatcaaagg attcgatggt atctgctcag catgggcagc 3300

aatgaaaaca tccattctat tcatttcagt ggacatgtgt tcactgtacg aaaaaaagag 3360

gagtataaaa tggcactgta caatctctat ccaggtgttt ttgagacagt ggaaatgtta 3420

ccatccaaag ctggaatttg gcgggtggaa tgccttattg gcgagcatct acatgctggg 3480

atgagcacac tttttctggt gtacagcaat aagtgtcaga ctcccctggg aatggcttct 3540

ggacacatta gagattttca gattacagct tcaggacaat atggacagtg ggccccaaag 3600

ctggccagac ttcattattc cggatcaatc aatgcctgga gcaccaagga gccctttct 3660

tggatcaagg tggatctgtt ggcaccaatg attattcacg gcatcaagac ccagggtgcc 3720

cgtcagaagt tctccagcct ctacatctct cagtttatca tcatgtatag tcttgatggg 3780

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cgtttgcacc caactcatta tagcattcgc agcactcttc gcatggagtt gatgggctgt 3960

gatttaaata gttgcagcat gccattggga atggagagta aagcaatatc agatgcacag 4020

attactgctt catcctactt taccaatatg tttgccacct ggtctccttc aaaagctcga 4080

cttcacctcc aagggaggag taatgcctgg agacctcagg tgaataatcc aaaagagtgg 4140

ctgcaagtgg acttccagaa gacaatgaaa gtcacaggag taactactca gggagtaaaa 4200

tctctgctta ccagcatgta tgtgaaggag ttcctcatct ccagcagtca agatggccat 4260

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ccccagagtt gggtgcacca gattgccctg aggatggagg ttctgggctg cgaggcacag 4440

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<210> 623
<211> 1514
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 623
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

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Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350
```

```
Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Ser Val Ala Gly Ala Pro Gly Ser Pro
    370                 375                 380

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr
385                 390                 395                 400

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
                405                 410                 415

Pro Ala Ser Ser Lys Lys His Pro Lys Thr Trp Val His Tyr Ile Ala
            420                 425                 430

Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu Ala Pro Asp
        435                 440                 445

Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile
    450                 455                 460

Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr
465                 470                 475                 480

Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro
                485                 490                 495

Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn
            500                 505                 510

Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val
        515                 520                 525

Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys
    530                 535                 540

Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val
545                 550                 555                 560

Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg
                565                 570                 575

Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu
            580                 585                 590

Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly
        595                 600                 605

Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe
    610                 615                 620
```

```
Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu
625                 630                 635                 640

Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser
                645                 650                 655

Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu
            660                 665                 670

Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly
        675                 680                 685

Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys
    690                 695                 700

His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly
705                 710                 715                 720

Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly
                725                 730                 735

Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys
            740                 745                 750

Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr
        755                 760                 765

Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu Pro
    770                 775                 780

Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His Gln Arg Glu
785                 790                 795                 800

Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp
                805                 810                 815

Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp
            820                 825                 830

Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His
        835                 840                 845

Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser
    850                 855                 860

Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln
865                 870                 875                 880

Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln
                885                 890                 895
```

```
Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro
            900                 905                 910

Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn
        915                 920                 925

Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu
    930                 935                 940

Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro
945                 950                 955                 960

Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro
                965                 970                 975

Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val
            980                 985                 990

Asp Leu Glu Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
        995                 1000                1005

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1010                1015                1020

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1025                1030                1035

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1040                1045                1050

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1055                1060                1065

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1070                1075                1080

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1085                1090                1095

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1100                1105                1110

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1115                1120                1125

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1130                1135                1140

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1145                1150                1155
```

```
Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1160                1165                1170

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1175                1180                1185

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1190                1195                1200

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1205                1210                1215

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1220                1225                1230

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1235                1240                1245

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1250                1255                1260

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1265                1270                1275

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1280                1285                1290

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1295                1300                1305

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1310                1315                1320

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1325                1330                1335

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1340                1345                1350

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1355                1360                1365

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1370                1375                1380

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1385                1390                1395

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1400                1405                1410
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Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1415 1420 1425

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1430 1435 1440

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1445 1450 1455

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1460 1465 1470

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1475 1480 1485

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1490 1495 1500

Ser Pro Ala Thr Gly His His His His His His
1505 1510

<210> 624
<211> 4545
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

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atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgcctctgt agcaggcgcg 1140

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ggacacatta gagattttca gattacagct tcaggacaat atggacagtg ggccccaaag 3600

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aagaagtggc agacttatcg aggaaattcc actggaacct taatggtctt ctttggcaat 3840

gtggattcat ctgggataaa acacaatatt tttaaccctc caattattgc tcgatacatc 3900

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<210> 625
<211> 1511
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile

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        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr
                405                 410                 415

Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
            420                 425                 430

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Ala Ser Ser Asp Arg Ser
        435                 440                 445

Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys
    450                 455                 460

Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr
465                 470                 475                 480

Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr
                485                 490                 495

Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser
            500                 505                 510

Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu
        515                 520                 525

Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro
    530                 535                 540

Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu
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545 550 555 560

Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser
565 570 575

Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro
580 585 590

Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile
595 600 605

Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn
610 615 620

Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro
625 630 635 640

Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met
645 650 655

His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys
660 665 670

Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr
675 680 685

Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met
690 695 700

Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val
705 710 715 720

Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn
725 730 735

Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser
740 745 750

Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile
755 760 765

Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe
770 775 780

Ser Gln Asn Pro Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg
785 790 795 800

Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
805 810 815

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu

820 825 830

Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile
835 840 845

Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His
850 855 860

Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys
865 870 875 880

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
885 890 895

Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
900 905 910

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser
915 920 925

Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln
930 935 940

Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
945 950 955 960

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp
965 970 975

Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu
980 985 990

Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val Cys His Thr
995 1000 1005

Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr Val Gln Glu
1010 1015 1020

Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys Ser Trp Tyr
1025 1030 1035

Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro Cys Asn Ile
1040 1045 1050

Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg Phe His Ala
1055 1060 1065

Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu Val Met Ala
1070 1075 1080

Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met Gly Ser Asn

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Glu Asn  Ile His Ser Ile His  Phe Ser Gly His Val  Phe Thr Val
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Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
    1115                1120                1125

Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    1130                1135                1140

Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    1145                1150                1155

Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    1160                1165                1170

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
    1175                1180                1185

Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
    1190                1195                1200

Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
    1205                1210                1215

Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
    1220                1225                1230

Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    1235                1240                1245

Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
    1250                1255                1260

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    1265                1270                1275

Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
    1280                1285                1290

Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1295                1300                1305

Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
    1310                1315                1320

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    1325                1330                1335

Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
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1340 1345 1350

Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg
1355 1360 1365

Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln
1370 1375 1380

Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser
1385 1390 1395

Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser
1400 1405 1410

Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val
1415 1420 1425

Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn
1430 1435 1440

Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro
1445 1450 1455

Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly
1460 1465 1470

Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu
1475 1480 1485

Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala
1490 1495 1500

Thr Gly His His His His His His
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<210> 626
<211> 4536
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

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gttgttttcc aggaatttac tgatggctcc tttactcagc ccttataccg tggagaacta 2700

aatgaacatt tgggactcct ggggccatat ataagagcag aagttgaaga taatatcatg 2760

gtaactttca gaaatcaggc ctctcgtccc tattccttct attctagcct tatttcttat 2820

gaggaagatc agaggcaagg agcagaacct agaaaaaact ttgtcaagcc taatgaaacc 2880

aaaacttact tttggaaagt gcaacatcat atggcaccca ctaaagatga gtttgactgc 2940

aaagcctggg cttatttctc tgatgttgac ctggaaaaag atgtgcactc aggcctgatt 3000

ggaccccttc tggtctgcca cactaacaca ctgaaccctg ctcatgggag acaagtgaca 3060

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gaaaatatgg aaagaaactg cagggctccc tgcaatatcc agatggaaga tcccactttt 3180

aaagagaatt atcgcttcca tgcaatcaat ggctacataa tggatacact acctggctta 3240

gtaatggctc aggatcaaag gattcgatgg tatctgctca gcatgggcag caatgaaaac 3300

atccattcta ttcatttcag tggacatgtg ttcactgtac gaaaaaaaga ggagtataaa 3360

atggcactgt acaatctcta tccaggtgtt tttgagacag tggaaatgtt accatccaaa 3420

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ccaactcatt atagcattcg cagcactctt cgcatggagt tgatgggctg tgatttaaat 3960

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tcatcctact ttaccaatat gtttgccacc tggtctcctt caaaagctcg acttcacctc 4080

caagggagga gtaatgcctg gagacctcag gtgaataatc caaaagagtg gctgcaagtg 4140

gacttccaga agacaatgaa agtcacagga gtaactactc agggagtaaa atctctgctt 4200

accagcatgt atgtgaagga gttcctcatc tccagcagtc aagatggcca tcagtggact 4260

ctctttttc agaatggcaa agtaaaggtt tttcagggaa atcaagactc cttcacacct 4320

gtggtgaact ctctagaccc accgttactg actcgctacc ttcgaattca cccccagagt 4380

tgggtgcacc agattgccct gaggatggag gttctgggct gcgaggcaca ggacctctac 4440

ggcgccggat cacctggggc cgaaacggcc gaacaaaaac tcatctcaga agaggatctg 4500

tcacctgcaa ccggtcatca tcaccatcac cattga 4536

<210> 627
<211> 1511
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 627
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

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His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gly Ala Pro Gly Ser Pro
        435                 440                 445

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr
    450                 455                 460

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
465                 470                 475                 480
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Pro Ala Ser Ser Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr
                485                 490                 495

Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser
            500                 505                 510

Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu
        515                 520                 525

Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro
    530                 535                 540

Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu
545                 550                 555                 560

Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser
                565                 570                 575

Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro
            580                 585                 590

Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile
        595                 600                 605

Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn
    610                 615                 620

Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro
625                 630                 635                 640

Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met
                645                 650                 655

His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys
            660                 665                 670

Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr
        675                 680                 685

Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met
    690                 695                 700

Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val
705                 710                 715                 720

Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn
                725                 730                 735

Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser
            740                 745                 750
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Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile
        755                 760                 765

Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe
    770                 775                 780

Ser Gln Asn Pro Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg
785                 790                 795                 800

Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
                805                 810                 815

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu
            820                 825                 830

Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile
        835                 840                 845

Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His
    850                 855                 860

Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys
865                 870                 875                 880

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
                885                 890                 895

Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
            900                 905                 910

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser
        915                 920                 925

Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln
    930                 935                 940

Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
945                 950                 955                 960

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp
                965                 970                 975

Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu
            980                 985                 990

Lys Asp Val His Ser Gly Leu Ile  Gly Pro Leu Leu Val  Cys His Thr
        995                 1000                1005

Asn Thr  Leu Asn Pro Ala His  Gly Arg Gln Val Thr  Val Gln Glu
    1010                1015                1020
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```
Phe Ala  Leu Phe Phe Thr Ile  Phe Asp Glu Thr Lys  Ser Trp Tyr
    1025                1030                1035

Phe Thr  Glu Asn Met Glu Arg  Asn Cys Arg Ala Pro  Cys Asn Ile
    1040                1045                1050

Gln Met  Glu Asp Pro Thr Phe  Lys Glu Asn Tyr Arg  Phe His Ala
    1055                1060                1065

Ile Asn  Gly Tyr Ile Met Asp  Thr Leu Pro Gly Leu  Val Met Ala
    1070                1075                1080

Gln Asp  Gln Arg Ile Arg Trp  Tyr Leu Leu Ser Met  Gly Ser Asn
    1085                1090                1095

Glu Asn  Ile His Ser Ile His  Phe Ser Gly His Val  Phe Thr Val
    1100                1105                1110

Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
    1115                1120                1125

Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    1130                1135                1140

Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    1145                1150                1155

Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    1160                1165                1170

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
    1175                1180                1185

Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
    1190                1195                1200

Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
    1205                1210                1215

Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
    1220                1225                1230

Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    1235                1240                1245

Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
    1250                1255                1260

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    1265                1270                1275
```

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Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
    1280                1285                1290

Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1295                1300                1305

Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
    1310                1315                1320

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    1325                1330                1335

Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
    1340                1345                1350

Ser Lys  Ala Arg Leu His Leu  Gln Gly Arg Ser Asn  Ala Trp Arg
    1355                1360                1365

Pro Gln  Val Asn Asn Pro Lys  Glu Trp Leu Gln Val  Asp Phe Gln
    1370                1375                1380

Lys Thr  Met Lys Val Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser
    1385                1390                1395

Leu Leu  Thr Ser Met Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser
    1400                1405                1410

Gln Asp  Gly His Gln Trp Thr  Leu Phe Phe Gln Asn  Gly Lys Val
    1415                1420                1425

Lys Val  Phe Gln Gly Asn Gln  Asp Ser Phe Thr Pro  Val Val Asn
    1430                1435                1440

Ser Leu  Asp Pro Pro Leu Leu  Thr Arg Tyr Leu Arg  Ile His Pro
    1445                1450                1455

Gln Ser  Trp Val His Gln Ile  Ala Leu Arg Met Glu  Val Leu Gly
    1460                1465                1470

Cys Glu  Ala Gln Asp Leu Tyr  Gly Ala Gly Ser Pro  Gly Ala Glu
    1475                1480                1485

Thr Ala  Glu Gln Lys Leu Ile  Ser Glu Glu Asp Leu  Ser Pro Ala
    1490                1495                1500

Thr Gly  His His His His His  His
    1505                1510
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<210> 628
<211> 4536
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 628
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ctcggtgagc tgcctgtgga cgcaagattt cctcctagag tgccaaaatc ttttccattc 120

aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

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gctattggcg cgccaggttc tcctgctggc tcccccacct caacagaaga ggggacaagc 1380

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tttgatgaga accgaagctg gtacctcaca gagaatatac aacgctttct ccccaatcca 1920

gctggagtgc agcttgagga tccagagttc caagcctcca acatcatgca cagcatcaat 1980

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gtaactttca gaaatcaggc ctctcgtccc tattccttct attctagcct tatttcttat 2820

gaggaagatc agaggcaagg agcagaacct agaaaaaact ttgtcaagcc taatgaaacc 2880

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cagacttatc gaggaaattc cactggaacc ttaatggtct tctttggcaa tgtggattca 3840

tctgggataa aacacaatat ttttaaccct ccaattattg ctcgatacat ccgtttgcac 3900

ccaactcatt atagcattcg cagcactctt cgcatggagt tgatgggctg tgatttaaat 3960

agttgcagca tgccattggg aatggagagt aaagcaatat cagatgcaca gattactgct 4020

tcatcctact ttaccaatat gtttgccacc tggtctcctt caaaagctcg acttcacctc 4080

caagggagga gtaatgcctg gagacctcag gtgaataatc caaaagagtg gctgcaagtg 4140

gacttccaga agacaatgaa agtcacagga gtaactactc agggagtaaa atctctgctt 4200

accagcatgt atgtgaagga gttcctcatc tccagcagtc aagatggcca tcagtggact 4260

ctctttttc agaatggcaa agtaaaggtt tttcagggaa atcaagactc cttcacacct 4320

gtggtgaact ctctagaccc accgttactg actcgctacc ttcgaattca cccccagagt 4380

tgggtgcacc agattgccct gaggatggag gttctgggct gcgaggcaca ggacctctac 4440

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tcacctgcaa ccggtcatca tcaccatcac cattga 4536

<210> 629
<211> 1514
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 629

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

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Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415
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Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Ala Pro
                485                 490                 495

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
            500                 505                 510

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
        515                 520                 525

Ser Glu Thr Pro Ala Ser Ser Leu Pro Lys Gly Val Lys His Leu Lys
    530                 535                 540

Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val
545                 550                 555                 560

Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg
                565                 570                 575

Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu
            580                 585                 590

Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly
        595                 600                 605

Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe
    610                 615                 620

Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu
625                 630                 635                 640

Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser
                645                 650                 655

Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu
            660                 665                 670

Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly
        675                 680                 685
```

```
Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys
    690                 695                 700

His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly
705                 710                 715                 720

Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly
                725                 730                 735

Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys
            740                 745                 750

Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr
        755                 760                 765

Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu Pro
    770                 775                 780

Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His Gln Arg Glu
785                 790                 795                 800

Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp
                805                 810                 815

Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp
            820                 825                 830

Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His
        835                 840                 845

Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser
    850                 855                 860

Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln
865                 870                 875                 880

Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln
                885                 890                 895

Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro
            900                 905                 910

Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn
        915                 920                 925

Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu
    930                 935                 940

Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro
945                 950                 955                 960
```

```
Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro
               965                 970                 975

Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val
           980                 985                 990

Asp Leu Glu Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
       995                1000                1005

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
   1010                1015                1020

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
   1025                1030                1035

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
   1040                1045                1050

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
   1055                1060                1065

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
   1070                1075                1080

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
   1085                1090                1095

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
   1100                1105                1110

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
   1115                1120                1125

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
   1130                1135                1140

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
   1145                1150                1155

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
   1160                1165                1170

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
   1175                1180                1185

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
   1190                1195                1200

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
   1205                1210                1215
```

```
Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1220                1225                1230

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1235                1240                1245

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1250                1255                1260

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1265                1270                1275

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1280                1285                1290

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1295                1300                1305

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1310                1315                1320

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1325                1330                1335

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1340                1345                1350

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1355                1360                1365

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1370                1375                1380

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1385                1390                1395

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1400                1405                1410

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1415                1420                1425

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1430                1435                1440

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1445                1450                1455

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1460                1465                1470
```

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1475 1480 1485

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1490 1495 1500

Ser Pro Ala Thr Gly His His His His His His
1505 1510

<210> 630
<211> 4545
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 630
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ctcggtgagc tgcctgtgga cgcaagattt cctcctagag tgccaaaatc ttttccattc 120

aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ctccctaaag gcgcgccagg ttctcctgct 1500

ggctccccca cctcaacaga agaggggaca agcgaaagcg ctacgcctga gagtggccct 1560

ggctctgagc cagccacctc cggctctgaa accctgcct cgagcttacc aaaaggtgta 1620

aaacatttga aggattttcc aattctgcca ggagaaatat tcaaatataa atggacagtg 1680

actgtagaag atgggccaac taaatcagat cctcggtgcc tgacccgcta ttactctagt 1740

ttcgttaata tggagagaga tctagcttca ggactcattg gccctctcct catctgctac 1800

aaagaatctg tagatcaaag aggaaaccag ataatgtcag acaagaggaa tgtcatcctg 1860

ttttctgtat ttgatgagaa ccgaagctgg tacctcacag agaatataca acgctttctc 1920

cccaatccag ctggagtgca gcttgaggat ccagagttcc aagcctccaa catcatgcac 1980

agcatcaatg gctatgtttt tgatagtttg cagttgtcag tttgtttgca tgaggtggca 2040

tactggtaca ttctaagcat tggagcacag actgacttcc tttctgtctt cttctctgga 2100

tataccttca aacacaaaat ggtctatgaa gacacactca ccctattccc attctcagga 2160

gaaactgtct tcatgtcgat ggaaaaccca ggtctatgga ttctggggtg ccacaactca 2220

gactttcgga acagaggcat gaccgcctta ctgaaggttt ctagttgtga caagaacact 2280

ggtgattatt acgaggacag ttatgaagat atttcagcat acttgctgag taaaaacaat 2340

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gttgaaatga agaaggaaga ttttgacatt tatgatgagg atgaaaatca gagcccccgc 2520

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gggatgagta gctccccaca tgttctaaga aacagggctc agagtggcag tgtccctcag 2640

ttcaagaaag ttgttttcca ggaatttact gatggctcct ttactcagcc cttataccgt 2700

ggagaactaa atgaacattt gggactcctg gggccatata taagagcaga agttgaagat 2760

aatatcatgg taactttcag aaatcaggcc tctcgtccct attccttcta ttctagcctt 2820

atttcttatg aggaagatca gaggcaagga gcagaaccta gaaaaaactt tgtcaagcct 2880

aatgaaacca aaacttactt ttggaaagtg caacatcata tggcacccac taaagatgag 2940

tttgactgca aagcctgggc ttatttctct gatgttgacc tggaaaaaga tgtgcactca 3000

ggcctgattg gaccccttct ggtctgccac actaacacac tgaaccctgc tcatgggaga 3060

caagtgacag tacaggaatt tgctctgttt ttcaccatct ttgatgagac caaaagctgg 3120

tacttcactg aaaatatgga aagaaactgc agggctccct gcaatatcca gatggaagat 3180

cccactttta aagagaatta tcgcttccat gcaatcaatg gctacataat ggatacacta 3240

cctggcttag taatggctca ggatcaaagg attcgatggt atctgctcag catgggcagc 3300

aatgaaaaca tccattctat tcatttcagt ggacatgtgt tcactgtacg aaaaaaagag 3360

gagtataaaa tggcactgta caatctctat ccaggtgttt ttgagacagt ggaaatgtta 3420

ccatccaaag ctggaatttg gcgggtggaa tgccttattg gcgagcatct acatgctggg 3480

atgagcacac tttttctggt gtacagcaat aagtgtcaga ctcccctggg aatggcttct 3540

ggacacatta gagattttca gattacagct tcaggacaat atggacagtg ggccccaaag 3600

ctggccagac ttcattattc cggatcaatc aatgcctgga gcaccaagga gccctttct 3660

tggatcaagg tggatctgtt ggcaccaatg attattcacg gcatcaagac ccagggtgcc 3720

cgtcagaagt tctccagcct ctacatctct cagtttatca tcatgtatag tcttgatggg 3780

aagaagtggc agacttatcg aggaaattcc actggaacct taatggtctt ctttggcaat 3840

gtggattcat ctgggataaa acacaatatt tttaaccctc caattattgc tcgatacatc 3900

cgtttgcacc caactcatta tagcattcgc agcactcttc gcatggagtt gatgggctgt 3960

gatttaaata gttgcagcat gccattggga atggagagta aagcaatatc agatgcacag 4020

attactgctt catcctactt taccaatatg tttgccacct ggtctccttc aaaagctcga 4080

cttcacctcc aagggaggag taatgcctgg agacctcagg tgaataatcc aaaagagtgg 4140

ctgcaagtgg acttccagaa gacaatgaaa gtcacaggag taactactca gggagtaaaa 4200

tctctgctta ccagcatgta tgtgaaggag ttcctcatct ccagcagtca agatggccat 4260

cagtggactc tctttttca gaatggcaaa gtaaaggttt ttcagggaaa tcaagactcc 4320

ttcacacctg tggtgaactc tctagaccca ccgttactga ctcgctacct tcgaattcac 4380

ccccagagtt gggtgcacca gattgccctg aggatggagg ttctgggctg cgaggcacag 4440

gacctctacg gcgccggatc acctggggcc gaaacggccg aacaaaaact catctcagaa 4500

gaggatctgt cacctgcaac cggtcatcat caccatcacc attga 4545

<210> 631
<211> 1511
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 631
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

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Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350
```

```
Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Gly Ala Pro Gly Ser Pro Ala Gly Ser Pro
        515                 520                 525

Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    530                 535                 540

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Ala Ser Ser
545                 550                 555                 560

Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser
                565                 570                 575

Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro
            580                 585                 590

Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile
        595                 600                 605

Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn
    610                 615                 620
```

```
Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro
625                 630                 635                 640

Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met
                645                 650                 655

His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys
            660                 665                 670

Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr
        675                 680                 685

Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met
    690                 695                 700

Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val
705                 710                 715                 720

Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn
                725                 730                 735

Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser
            740                 745                 750

Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile
        755                 760                 765

Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe
    770                 775                 780

Ser Gln Asn Pro Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg
785                 790                 795                 800

Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
                805                 810                 815

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu
            820                 825                 830

Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile
        835                 840                 845

Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His
    850                 855                 860

Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys
865                 870                 875                 880

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
                885                 890                 895
```

```
Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
            900                 905                 910

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser
        915                 920                 925

Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln
    930                 935                 940

Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
945                 950                 955                 960

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp
                965                 970                 975

Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu
            980                 985                 990

Lys Asp Val His Ser Gly Leu Ile  Gly Pro Leu Leu Val  Cys His Thr
        995                 1000                1005

Asn Thr  Leu Asn Pro Ala His  Gly Arg Gln Val Thr  Val Gln Glu
    1010                1015                1020

Phe Ala  Leu Phe Phe Thr Ile  Phe Asp Glu Thr Lys  Ser Trp Tyr
    1025                1030                1035

Phe Thr  Glu Asn Met Glu Arg  Asn Cys Arg Ala Pro  Cys Asn Ile
    1040                1045                1050

Gln Met  Glu Asp Pro Thr Phe  Lys Glu Asn Tyr Arg  Phe His Ala
    1055                1060                1065

Ile Asn  Gly Tyr Ile Met Asp  Thr Leu Pro Gly Leu  Val Met Ala
    1070                1075                1080

Gln Asp  Gln Arg Ile Arg Trp  Tyr Leu Leu Ser Met  Gly Ser Asn
    1085                1090                1095

Glu Asn  Ile His Ser Ile His  Phe Ser Gly His Val  Phe Thr Val
    1100                1105                1110

Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
    1115                1120                1125

Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    1130                1135                1140

Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    1145                1150                1155
```

```
Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    1160                1165                1170

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
    1175                1180                1185

Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
    1190                1195                1200

Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
    1205                1210                1215

Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
    1220                1225                1230

Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    1235                1240                1245

Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
    1250                1255                1260

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    1265                1270                1275

Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
    1280                1285                1290

Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1295                1300                1305

Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
    1310                1315                1320

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    1325                1330                1335

Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
    1340                1345                1350

Ser Lys  Ala Arg Leu His Leu  Gln Gly Arg Ser Asn  Ala Trp Arg
    1355                1360                1365

Pro Gln  Val Asn Asn Pro Lys  Glu Trp Leu Gln Val  Asp Phe Gln
    1370                1375                1380

Lys Thr  Met Lys Val Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser
    1385                1390                1395

Leu Leu  Thr Ser Met Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser
    1400                1405                1410
```

Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val
1415 1420 1425

Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn
1430 1435 1440

Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro
1445 1450 1455

Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly
1460 1465 1470

Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu
1475 1480 1485

Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala
1490 1495 1500

Thr Gly His His His His His His
1505 1510

<210> 632
<211> 4536
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 632
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
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His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
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Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
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Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
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Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
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Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
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Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile

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Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
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Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
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Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
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Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
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Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
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Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
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Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
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Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
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Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
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Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
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Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
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His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
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Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
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Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
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Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
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Arg Phe Leu Pro Asn Pro Ala Gly Ala Pro Gly Ser Pro Ala Gly Ser
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Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
610 615 620

Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Ala Ser
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Ser Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met
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His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys
660 665 670

Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr
675 680 685

Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met
690 695 700

Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val
705 710 715 720

Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn
725 730 735

Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser
740 745 750

Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile
755 760 765

Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe
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Ser Gln Asn Pro Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg
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Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu

820 825 830

Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile
835 840 845

Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His
850 855 860

Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys
865 870 875 880

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
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Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
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Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser
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Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln
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Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
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Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp
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Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val Cys His Thr
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Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
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Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
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Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
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Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
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Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
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Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
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Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
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Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
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Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
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Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn
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Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly
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ctctttttc agaatggcaa agtaaaggtt tttcagggaa atcaagactc cttcacacct 4320

gtggtgaact ctctagaccc accgttactg actcgctacc ttcgaattca cccccagagt 4380

tgggtgcacc agattgccct gaggatggag gttctgggct gcgaggcaca ggacctctac 4440

ggcgccggat cacctggggc cgaaacggcc gaacaaaaac tcatctcaga agaggatctg 4500

tcacctgcaa ccggtcatca tcaccatcac cattga 4536

<210> 635
<211> 1511
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 635
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

```
His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480
```

```
Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Gly Ala Pro Gly Ser Pro Ala
705                 710                 715                 720

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
                725                 730                 735

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
            740                 745                 750
```

```
Ala Ser Ser Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile
        755                 760                 765

Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe
    770                 775                 780

Ser Gln Asn Pro Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg
785                 790                 795                 800

Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
                805                 810                 815

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu
            820                 825                 830

Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile
        835                 840                 845

Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His
    850                 855                 860

Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys
865                 870                 875                 880

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
                885                 890                 895

Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
            900                 905                 910

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser
        915                 920                 925

Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln
    930                 935                 940

Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
945                 950                 955                 960

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp
                965                 970                 975

Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu
            980                 985                 990

Lys Asp Val His Ser Gly Leu Ile  Gly Pro Leu Leu Val  Cys His Thr
        995                 1000                1005

Asn Thr  Leu Asn Pro Ala His  Gly Arg Gln Val Thr  Val Gln Glu
    1010                1015                1020
```

```
Phe Ala  Leu Phe Phe Thr Ile  Phe Asp Glu Thr Lys  Ser Trp Tyr
    1025                1030                1035

Phe Thr  Glu Asn Met Glu Arg  Asn Cys Arg Ala Pro  Cys Asn Ile
    1040                1045                1050

Gln Met  Glu Asp Pro Thr Phe  Lys Glu Asn Tyr Arg  Phe His Ala
    1055                1060                1065

Ile Asn  Gly Tyr Ile Met Asp  Thr Leu Pro Gly Leu  Val Met Ala
    1070                1075                1080

Gln Asp  Gln Arg Ile Arg Trp  Tyr Leu Leu Ser Met  Gly Ser Asn
    1085                1090                1095

Glu Asn  Ile His Ser Ile His  Phe Ser Gly His Val  Phe Thr Val
    1100                1105                1110

Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
    1115                1120                1125

Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    1130                1135                1140

Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    1145                1150                1155

Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    1160                1165                1170

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
    1175                1180                1185

Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
    1190                1195                1200

Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
    1205                1210                1215

Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
    1220                1225                1230

Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    1235                1240                1245

Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
    1250                1255                1260

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    1265                1270                1275
```

Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro Pro Ile Ile
1280 1285 1290

Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser Ile Arg Ser
1295 1300 1305

Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn Ser Cys Ser
1310 1315 1320

Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp Ala Gln Ile
1325 1330 1335

Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp Ser Pro
1340 1345 1350

Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg
1355 1360 1365

Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln
1370 1375 1380

Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser
1385 1390 1395

Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser
1400 1405 1410

Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val
1415 1420 1425

Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn
1430 1435 1440

Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro
1445 1450 1455

Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly
1460 1465 1470

Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu
1475 1480 1485

Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala
1490 1495 1500

Thr Gly His His His His His His
1505 1510

<210> 636
<211> 4536
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 636
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ctcggtgagc tgcctgtgga cgcaagattt cctcctagag tgccaaaatc ttttccattc 120

aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

agagatctag cttcaggact cattggccct ctcctcatct gctacaaaga atctgtagat 1680

caaagaggaa accagataat gtcagacaag aggaatgtca tcctgttttc tgtatttgat 1740

gagaaccgaa gctggtacct cacagagaat atacaacgct ttctccccaa tccagctgga 1800

gtgcagcttg aggatccaga gttccaagcc tccaacatca tgcacagcat caatggctat 1860

gtttttgata gtttgcagtt gtcagtttgt ttgcatgagg tggcatactg gtacattcta 1920

agcattggag cacagactga cttccttttct gtcttcttct ctggatatac cttcaaacac 1980

aaaatggtct atgaagacac actcacccta ttcccattct caggagaaac tgtcttcatg 2040

tcgatggaaa acccaggtct atggattctg gggtgccaca actcagactt tcggaacaga 2100

ggcatgaccg ccttactgaa ggtttctagt tgtgacaagg gcgcgccagg ttctcctgct 2160

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ggctctgagc cagccacctc cggctctgaa acccctgcct cgagcaacac tggtgattat 2280

tacgaggaca gttatgaaga tatttcagca tacttgctga gtaaaaacaa tgccattgaa 2340

ccaagaagct tctctcaaaa cccaccagtc ttgaaacgcc atcaacggga aataactcgt 2400

actactcttc agtcagatca agaggaaatc gattatgatg ataccatatc agttgaaatg 2460

aagaaggaag attttgacat ttatgatgag gatgaaaatc agagcccccg cagctttcaa 2520

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gttgttttcc aggaatttac tgatggctcc tttactcagc ccttataccg tggagaacta 2700

aatgaacatt tgggactcct ggggccatat ataagagcag aagttgaaga taatatcatg 2760

gtaactttca gaaatcaggc ctctcgtccc tattccttct attctagcct tatttcttat 2820

gaggaagatc agaggcaagg agcagaacct agaaaaaact ttgtcaagcc taatgaaacc 2880

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aaagcctggg cttatttctc tgatgttgac ctggaaaaag atgtgcactc aggcctgatt 3000

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aaagagaatt atcgcttcca tgcaatcaat ggctacataa tggatacact acctggctta 3240

gtaatggctc aggatcaaag gattcgatgg tatctgctca gcatgggcag caatgaaaac 3300

atccattcta ttcatttcag tggacatgtg ttcactgtac gaaaaaaaga ggagtataaa 3360

atggcactgt acaatctcta tccaggtgtt tttgagacag tggaaatgtt accatccaaa 3420

gctggaattt ggcgggtgga atgccttatt ggcgagcatc tacatgctgg gatgagcaca 3480

ctttttctgg tgtacagcaa taagtgtcag actcccctgg gaatggcttc tggacacatt 3540

agagattttc agattacagc ttcaggacaa tatggacagt gggccccaaa gctggccaga 3600

cttcattatt ccggatcaat caatgcctgg agcaccaagg agccctttc ttggatcaag 3660

gtggatctgt tggcaccaat gattattcac ggcatcaaga cccagggtgc ccgtcagaag 3720

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cagacttatc gaggaaattc cactggaacc ttaatggtct tctttggcaa tgtggattca 3840

tctgggataa aacacaatat ttttaaccct ccaattattg ctcgatacat ccgtttgcac 3900

ccaactcatt atagcattcg cagcactctt cgcatggagt tgatgggctg tgatttaaat 3960

agttgcagca tgccattggg aatggagagt aaagcaatat cagatgcaca gattactgct 4020

tcatcctact ttaccaatat gtttgccacc tggtctcctt caaaagctcg acttcacctc 4080

caagggagga gtaatgcctg gagacctcag gtgaataatc caaaagagtg gctgcaagtg 4140

gacttccaga agacaatgaa agtcacagga gtaactactc agggagtaaa atctctgctt 4200

accagcatgt atgtgaagga gttcctcatc tccagcagtc aagatggcca tcagtggact 4260

ctctttttc agaatggcaa agtaaaggtt tttcagggaa atcaagactc cttcacacct 4320

gtggtgaact ctctagaccc accgttactg actcgctacc ttcgaattca cccccagagt 4380

tgggtgcacc agattgccct gaggatggag gttctgggct gcgaggcaca ggacctctac 4440

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tcacctgcaa ccggtcatca tcaccatcac cattga 4536

<210> 637
<211> 1511
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 637

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

```
Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415
```

```
Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685
```

```
Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Ser Pro Ala
            740                 745                 750

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
        755                 760                 765

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
    770                 775                 780

Ala Ser Ser Pro Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg
785                 790                 795                 800

Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
                805                 810                 815

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu
            820                 825                 830

Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile
        835                 840                 845

Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His
    850                 855                 860

Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys
865                 870                 875                 880

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
                885                 890                 895

Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
            900                 905                 910

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser
        915                 920                 925

Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln
    930                 935                 940

Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
945                 950                 955                 960
```

```
Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp
                965                 970                 975

Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu
            980                 985                 990

Lys Asp Val His Ser Gly Leu Ile  Gly Pro Leu Leu Val  Cys His Thr
        995                 1000                1005

Asn Thr  Leu Asn Pro Ala His  Gly Arg Gln Val Thr  Val Gln Glu
    1010                1015                1020

Phe Ala  Leu Phe Phe Thr Ile  Phe Asp Glu Thr Lys  Ser Trp Tyr
    1025                1030                1035

Phe Thr  Glu Asn Met Glu Arg  Asn Cys Arg Ala Pro  Cys Asn Ile
    1040                1045                1050

Gln Met  Glu Asp Pro Thr Phe  Lys Glu Asn Tyr Arg  Phe His Ala
    1055                1060                1065

Ile Asn  Gly Tyr Ile Met Asp  Thr Leu Pro Gly Leu  Val Met Ala
    1070                1075                1080

Gln Asp  Gln Arg Ile Arg Trp  Tyr Leu Leu Ser Met  Gly Ser Asn
    1085                1090                1095

Glu Asn  Ile His Ser Ile His  Phe Ser Gly His Val  Phe Thr Val
    1100                1105                1110

Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
    1115                1120                1125

Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    1130                1135                1140

Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    1145                1150                1155

Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    1160                1165                1170

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
    1175                1180                1185

Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
    1190                1195                1200

Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
    1205                1210                1215
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Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
    1220                1225                1230

Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    1235                1240                1245

Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
    1250                1255                1260

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    1265                1270                1275

Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
    1280                1285                1290

Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1295                1300                1305

Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
    1310                1315                1320

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    1325                1330                1335

Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
    1340                1345                1350

Ser Lys  Ala Arg Leu His Leu  Gln Gly Arg Ser Asn  Ala Trp Arg
    1355                1360                1365

Pro Gln  Val Asn Asn Pro Lys  Glu Trp Leu Gln Val  Asp Phe Gln
    1370                1375                1380

Lys Thr  Met Lys Val Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser
    1385                1390                1395

Leu Leu  Thr Ser Met Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser
    1400                1405                1410

Gln Asp  Gly His Gln Trp Thr  Leu Phe Phe Gln Asn  Gly Lys Val
    1415                1420                1425

Lys Val  Phe Gln Gly Asn Gln  Asp Ser Phe Thr Pro  Val Val Asn
    1430                1435                1440

Ser Leu  Asp Pro Pro Leu Leu  Thr Arg Tyr Leu Arg  Ile His Pro
    1445                1450                1455

Gln Ser  Trp Val His Gln Ile  Ala Leu Arg Met Glu  Val Leu Gly
    1460                1465                1470
```

Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu
1475 1480 1485

Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala
1490 1495 1500

Thr Gly His His His His His His
1505 1510

<210> 638
<211> 4536
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

agagatctag cttcaggact cattggccct ctcctcatct gctacaaaga atctgtagat 1680

caaagaggaa accagataat gtcagacaag aggaatgtca tcctgttttc tgtatttgat 1740

gagaaccgaa gctggtacct cacagagaat atacaacgct ttctccccaa tccagctgga 1800

gtgcagcttg aggatccaga gttccaagcc tccaacatca tgcacagcat caatggctat 1860

gtttttgata gtttgcagtt gtcagtttgt ttgcatgagg tggcatactg gtacattcta 1920

agcattggag cacagactga cttccttttct gtcttcttct ctggatatac cttcaaacac 1980

aaaatggtct atgaagacac actcacccta ttcccattct caggagaaac tgtcttcatg 2040

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ggcatgaccg ccttactgaa ggtttctagt tgtgacaaga acactggtga ttattacgag 2160

gacagttatg aagatatttc agcatacttg ctgagtaaaa acaatgccat tgaaccaaga 2220

agcttctctc aaaacggcgc gccaggttct cctgctggct cccccacctc aacagaagag 2280

gggacaagcg aaagcgctac gcctgagagt ggccctggct ctgagccagc cacctccggc 2340

tctgaaaccc ctgcctcgag cccaccagtc ttgaaacgcc atcaacggga aataactcgt 2400

actactcttc agtcagatca agaggaaatc gattatgatg ataccatatc agttgaaatg 2460

aagaaggaag attttgacat ttatgatgag gatgaaaatc agagcccccg cagctttcaa 2520

aagaaaacac gacactattt tattgctgca gtggagaggc tctgggatta tgggatgagt 2580

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gttgttttcc aggaatttac tgatggctcc tttactcagc ccttataccg tggagaacta 2700

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gtaactttca gaaatcaggc ctctcgtccc tattccttct attctagcct tatttcttat 2820

gaggaagatc agaggcaagg agcagaacct agaaaaaact ttgtcaagcc taatgaaacc 2880

aaaacttact tttggaaagt gcaacatcat atggcaccca ctaaagatga gtttgactgc 2940

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ggaccccttc tggtctgcca cactaacaca ctgaaccctg ctcatgggag acaagtgaca 3060

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gtaatggctc aggatcaaag gattcgatgg tatctgctca gcatgggcag caatgaaaac 3300

atccattcta ttcatttcag tggacatgtg ttcactgtac gaaaaaaaga ggagtataaa 3360

atggcactgt acaatctcta tccaggtgtt tttgagacag tggaaatgtt accatccaaa 3420

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cagacttatc gaggaaattc cactggaacc ttaatggtct tctttggcaa tgtggattca 3840

tctgggataa aacacaatat ttttaaccct ccaattattg ctcgatacat ccgtttgcac 3900

ccaactcatt atagcattcg cagcactctt cgcatggagt tgatgggctg tgatttaaat 3960

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tcatcctact ttaccaatat gtttgccacc tggtctcctt caaaagctcg acttcacctc 4080

caagggagga gtaatgcctg gagacctcag gtgaataatc caaaagagtg gctgcaagtg 4140

gacttccaga agacaatgaa agtcacagga gtaactactc agggagtaaa atctctgctt 4200

accagcatgt atgtgaagga gttcctcatc tccagcagtc aagatggcca tcagtggact 4260

ctctttttc agaatggcaa agtaaaggtt tttcagggaa atcaagactc cttcacacct 4320

gtggtgaact ctctagaccc accgttactg actcgctacc ttcgaattca cccccagagt 4380

tgggtgcacc agattgccct gaggatggag gttctgggct gcgaggcaca ggacctctac 4440

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tcacctgcaa ccggtcatca tcaccatcac cattga 4536

<210> 639
<211> 1511
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

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Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350
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Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620
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Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Ser Pro Ala
            740                 745                 750

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
        755                 760                 765

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
    770                 775                 780

Ala Ser Ser Pro Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg
785                 790                 795                 800

Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
                805                 810                 815

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu
            820                 825                 830

Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile
        835                 840                 845

Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His
    850                 855                 860

Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys
865                 870                 875                 880

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
                885                 890                 895
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Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
            900                 905                 910

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser
        915                 920                 925

Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln
    930                 935                 940

Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
945                 950                 955                 960

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp
                965                 970                 975

Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu
            980                 985                 990

Lys Asp Val His Ser Gly Leu Ile  Gly Pro Leu Leu Val  Cys His Thr
        995                 1000                1005

Asn Thr  Leu Asn Pro Ala His  Gly Arg Gln Val Thr  Val Gln Glu
    1010                1015                1020

Phe Ala  Leu Phe Phe Thr Ile  Phe Asp Glu Thr Lys  Ser Trp Tyr
    1025                1030                1035

Phe Thr  Glu Asn Met Glu Arg  Asn Cys Arg Ala Pro  Cys Asn Ile
    1040                1045                1050

Gln Met  Glu Asp Pro Thr Phe  Lys Glu Asn Tyr Arg  Phe His Ala
    1055                1060                1065

Ile Asn  Gly Tyr Ile Met Asp  Thr Leu Pro Gly Leu  Val Met Ala
    1070                1075                1080

Gln Asp  Gln Arg Ile Arg Trp  Tyr Leu Leu Ser Met  Gly Ser Asn
    1085                1090                1095

Glu Asn  Ile His Ser Ile His  Phe Ser Gly His Val  Phe Thr Val
    1100                1105                1110

Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
    1115                1120                1125

Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    1130                1135                1140

Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    1145                1150                1155
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```
Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    1160                1165                1170

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
    1175                1180                1185

Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
    1190                1195                1200

Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
    1205                1210                1215

Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
    1220                1225                1230

Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    1235                1240                1245

Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
    1250                1255                1260

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    1265                1270                1275

Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
    1280                1285                1290

Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1295                1300                1305

Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
    1310                1315                1320

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    1325                1330                1335

Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
    1340                1345                1350

Ser Lys  Ala Arg Leu His Leu  Gln Gly Arg Ser Asn  Ala Trp Arg
    1355                1360                1365

Pro Gln  Val Asn Asn Pro Lys  Glu Trp Leu Gln Val  Asp Phe Gln
    1370                1375                1380

Lys Thr  Met Lys Val Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser
    1385                1390                1395

Leu Leu  Thr Ser Met Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser
    1400                1405                1410
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Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val
1415 1420 1425

Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn
1430 1435 1440

Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro
1445 1450 1455

Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly
1460 1465 1470

Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu
1475 1480 1485

Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala
1490 1495 1500

Thr Gly His His His His His His
1505 1510

<210> 640
<211> 4536
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

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gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

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<211> 1763
<212> PRT
<213> Artificial Sequence

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile

275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp

545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Thr Ser Glu
740 745 750

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
755 760 765

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
770 775 780

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
785 790 795 800

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
805 810 815

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu

820 825 830

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
835 840 845

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
850 855 860

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
865 870 875 880

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
885 890 895

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
900 905 910

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
915 920 925

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
930 935 940

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
945 950 955 960

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
965 970 975

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
980 985 990

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
995 1000 1005

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1010 1015 1020

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Ala Ser
1025 1030 1035

Ser Pro Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr
1040 1045 1050

Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
1055 1060 1065

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp
1070 1075 1080

Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr

1085 1090 1095

Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser
1100 1105 1110

Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro
1115 1120 1125

Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe
1130 1135 1140

Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
1145 1150 1155

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val
1160 1165 1170

Thr Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser
1175 1180 1185

Leu Ile Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg
1190 1195 1200

Lys Asn Phe Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys
1205 1210 1215

Val Gln His His Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys
1220 1225 1230

Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu Lys Asp Val His
1235 1240 1245

Ser Gly Leu Ile Gly Pro Leu Leu Val Cys His Thr Asn Thr Leu
1250 1255 1260

Asn Pro Ala His Gly Arg Gln Val Thr Val Gln Glu Phe Ala Leu
1265 1270 1275

Phe Phe Thr Ile Phe Asp Glu Thr Lys Ser Trp Tyr Phe Thr Glu
1280 1285 1290

Asn Met Glu Arg Asn Cys Arg Ala Pro Cys Asn Ile Gln Met Glu
1295 1300 1305

Asp Pro Thr Phe Lys Glu Asn Tyr Arg Phe His Ala Ile Asn Gly
1310 1315 1320

Tyr Ile Met Asp Thr Leu Pro Gly Leu Val Met Ala Gln Asp Gln
1325 1330 1335

Arg Ile Arg Trp Tyr Leu Leu Ser Met Gly Ser Asn Glu Asn Ile

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His Ser  Ile His Phe Ser Gly  His Val Phe Thr Val  Arg Lys Lys
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Glu Glu  Tyr Lys Met Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe
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Glu Thr  Val Glu Met Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val
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Glu Cys  Leu Ile Gly Glu His  Leu His Ala Gly Met  Ser Thr Leu
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Phe Leu  Val Tyr Ser Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala
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Ser Gly  His Ile Arg Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr
    1430                1435                1440

Gly Gln  Trp Ala Pro Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser
    1445                1450                1455

Ile Asn  Ala Trp Ser Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val
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Asp Leu  Leu Ala Pro Met Ile  Ile His Gly Ile Lys  Thr Gln Gly
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Ala Arg  Gln Lys Phe Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile
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Met Tyr  Ser Leu Asp Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn
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Ser Thr  Gly Thr Leu Met Val  Phe Phe Gly Asn Val  Asp Ser Ser
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Gly Ile  Lys His Asn Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr
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Ile Arg  Leu His Pro Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg
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Met Glu  Leu Met Gly Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu
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Gly Met  Glu Ser Lys Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser
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Ser Tyr  Phe Thr Asn Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala
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Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg Pro Gln Val
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Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln Lys Thr Met
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Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser Leu Leu Thr
1640 1645 1650

Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly
1655 1660 1665

His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val Lys Val Phe
1670 1675 1680

Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn Ser Leu Asp
1685 1690 1695

Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro Gln Ser Trp
1700 1705 1710

Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly Cys Glu Ala
1715 1720 1725

Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu Thr Ala Glu
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Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala Thr Gly His
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<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 643
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

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Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380
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Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655
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Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Thr Ser Glu
            740                 745                 750

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
        755                 760                 765

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
    770                 775                 780

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
785                 790                 795                 800

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
                805                 810                 815

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
            820                 825                 830

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
        835                 840                 845

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
    850                 855                 860

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
865                 870                 875                 880

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
                885                 890                 895

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
            900                 905                 910

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        915                 920                 925
```

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Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
    930                 935                 940

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
945                 950                 955                 960

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
                965                 970                 975

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
            980                 985                 990

Glu Pro Ser Glu Gly Ser Ala Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
        995                 1000                1005

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1010                1015                1020

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Ala Ser
    1025                1030                1035

Ser Pro  Pro Val Leu Lys Arg  His Gln Ala Glu Ile  Thr Arg Thr
    1040                1045                1050

Thr Leu  Gln Ser Asp Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile
    1055                1060                1065

Ser Val  Glu Met Lys Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp
    1070                1075                1080

Glu Asn  Gln Ser Pro Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr
    1085                1090                1095

Phe Ile  Ala Ala Val Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser
    1100                1105                1110

Ser Pro  His Val Leu Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro
    1115                1120                1125

Gln Phe  Lys Lys Val Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe
    1130                1135                1140

Thr Gln  Pro Leu Tyr Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu
    1145                1150                1155

Leu Gly  Pro Tyr Ile Arg Ala  Glu Val Glu Asp Asn  Ile Met Val
    1160                1165                1170

Thr Phe  Arg Asn Gln Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser
    1175                1180                1185
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```
Leu Ile  Ser Tyr Glu Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg
    1190                 1195                 1200

Lys Asn  Phe Val Lys Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys
    1205                 1210                 1215

Val Gln  His His Met Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys
    1220                 1225                 1230

Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu Glu Lys  Asp Val His
    1235                 1240                 1245

Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys His Thr  Asn Thr Leu
    1250                 1255                 1260

Asn Pro  Ala His Gly Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu
    1265                 1270                 1275

Phe Phe  Thr Ile Phe Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu
    1280                 1285                 1290

Asn Met  Glu Arg Asn Cys Arg  Ala Pro Cys Asn Ile  Gln Met Glu
    1295                 1300                 1305

Asp Pro  Thr Phe Lys Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly
    1310                 1315                 1320

Tyr Ile  Met Asp Thr Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln
    1325                 1330                 1335

Arg Ile  Arg Trp Tyr Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile
    1340                 1345                 1350

His Ser  Ile His Phe Ser Gly  His Val Phe Thr Val  Arg Lys Lys
    1355                 1360                 1365

Glu Glu  Tyr Lys Met Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe
    1370                 1375                 1380

Glu Thr  Val Glu Met Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val
    1385                 1390                 1395

Glu Cys  Leu Ile Gly Glu His  Leu His Ala Gly Met  Ser Thr Leu
    1400                 1405                 1410

Phe Leu  Val Tyr Ser Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala
    1415                 1420                 1425

Ser Gly  His Ile Arg Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr
    1430                 1435                 1440
```

```
Gly Gln  Trp Ala Pro Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser
    1445                 1450                 1455

Ile Asn  Ala Trp Ser Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val
    1460                 1465                 1470

Asp Leu  Leu Ala Pro Met Ile  Ile His Gly Ile Lys  Thr Gln Gly
    1475                 1480                 1485

Ala Arg  Gln Lys Phe Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile
    1490                 1495                 1500

Met Tyr  Ser Leu Asp Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn
    1505                 1510                 1515

Ser Thr  Gly Thr Leu Met Val  Phe Phe Gly Asn Val  Asp Ser Ser
    1520                 1525                 1530

Gly Ile  Lys His Asn Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr
    1535                 1540                 1545

Ile Arg  Leu His Pro Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg
    1550                 1555                 1560

Met Glu  Leu Met Gly Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu
    1565                 1570                 1575

Gly Met  Glu Ser Lys Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser
    1580                 1585                 1590

Ser Tyr  Phe Thr Asn Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala
    1595                 1600                 1605

Arg Leu  His Leu Gln Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val
    1610                 1615                 1620

Asn Asn  Pro Lys Glu Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met
    1625                 1630                 1635

Lys Val  Thr Gly Val Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr
    1640                 1645                 1650

Ser Met  Tyr Val Lys Glu Phe  Leu Ile Ser Ser Ser  Gln Asp Gly
    1655                 1660                 1665

His Gln  Trp Thr Leu Phe Phe  Gln Asn Gly Lys Val  Lys Val Phe
    1670                 1675                 1680

Gln Gly  Asn Gln Asp Ser Phe  Thr Pro Val Val Asn  Ser Leu Asp
    1685                 1690                 1695
```

Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro Gln Ser Trp
1700 1705 1710

Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly Cys Glu Ala
1715 1720 1725

Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu Thr Ala Glu
1730 1735 1740

Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala Thr Gly His
1745 1750 1755

His His His His His
1760

<210> 644
<211> 5292
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

agagatctag cttcaggact cattggccct ctcctcatct gctacaaaga atctgtagat 1680

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gaacctagaa aaaactttgt caagcctaat gaaaccaaaa cttacttttg gaaagtgcaa 3660

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accatctttg atgagaccaa aagctggtac ttcactgaaa atatggaaag aaactgcagg 3900

gctccctgca atatccagat ggaagatccc acttttaaag agaattatcg cttccatgca 3960

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ttactgactc gctaccttcg aattcacccc cagagttggg tgcaccagat tgccctgagg 5160

atggaggttc tgggctgcga ggcacaggac ctctacggcg ccggatcacc tggggccgaa 5220

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catcaccatt ga 5292

<210> 645
<211> 1514
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 645

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

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His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480
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```
Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750
```

```
Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Gly Ala Pro
            820                 825                 830

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
        835                 840                 845

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
    850                 855                 860

Ser Glu Thr Pro Ala Ser Ser Arg Ala Gln Ser Gly Ser Val Pro Gln
865                 870                 875                 880

Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln
                885                 890                 895

Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro
            900                 905                 910

Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn
        915                 920                 925

Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu
    930                 935                 940

Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro
945                 950                 955                 960

Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro
                965                 970                 975

Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val
            980                 985                 990

Asp Leu Glu Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
        995                 1000                1005

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1010                1015                1020
```

```
Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1025                1030                1035

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1040                1045                1050

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1055                1060                1065

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1070                1075                1080

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1085                1090                1095

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1100                1105                1110

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1115                1120                1125

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1130                1135                1140

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1145                1150                1155

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1160                1165                1170

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1175                1180                1185

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1190                1195                1200

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1205                1210                1215

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1220                1225                1230

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1235                1240                1245

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1250                1255                1260

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1265                1270                1275
```

```
Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1280                1285                1290

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1295                1300                1305

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1310                1315                1320

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1325                1330                1335

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1340                1345                1350

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1355                1360                1365

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1370                1375                1380

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1385                1390                1395

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1400                1405                1410

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1415                1420                1425

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1430                1435                1440

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1445                1450                1455

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1460                1465                1470

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1475                1480                1485

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
    1490                1495                1500

Ser Pro  Ala Thr Gly His His  His His His His His
    1505                1510
```

<210> 646
<211> 4545
<212> DNA

<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 646
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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

agagatctag cttcaggact cattggccct ctcctcatct gctacaaaga atctgtagat 1680

caaagaggaa accagataat gtcagacaag aggaatgtca tcctgttttc tgtatttgat 1740

gagaaccgaa gctggtacct cacagagaat atacaacgct ttctccccaa tccagctgga 1800

gtgcagcttg aggatccaga gttccaagcc tccaacatca tgcacagcat caatggctat 1860

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ggacacatta gagattttca gattacagct tcaggacaat atggacagtg ggccccaaag 3600
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cgtcagaagt tctccagcct ctacatctct cagtttatca tcatgtatag tcttgatggg 3780
aagaagtggc agacttatcg aggaaattcc actggaacct taatggtctt ctttggcaat 3840
gtggattcat ctgggataaa acacaatatt tttaaccctc caattattgc tcgatacatc 3900

cgtttgcacc caactcatta tagcattcgc agcactcttc gcatggagtt gatgggctgt 3960

gatttaaata gttgcagcat gccattggga atggagagta aagcaatatc agatgcacag 4020

attactgctt catcctactt taccaatatg tttgccacct ggtctccttc aaaagctcga 4080

cttcacctcc aagggaggag taatgcctgg agacctcagg tgaataatcc aaaagagtgg 4140

ctgcaagtgg acttccagaa gacaatgaaa gtcacaggag taactactca gggagtaaaa 4200

tctctgctta ccagcatgta tgtgaaggag ttcctcatct ccagcagtca agatggccat 4260

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ccccagagtt gggtgcacca gattgccctg aggatggagg ttctgggctg cgaggcacag 4440

gacctctacg gcgccggatc acctggggcc gaaacggccg aacaaaaact catctcagaa 4500

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<210> 647
<211> 1511
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 647
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

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Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415
```

```
Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685
```

```
Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Gly Ala Pro Gly Ser Pro Ala Gly Ser Pro
            900                 905                 910

Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
        915                 920                 925

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Ala Ser Ser
    930                 935                 940

Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
945                 950                 955                 960
```

```
Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp
                965                 970                 975

Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu
            980                 985                 990

Lys Asp Val His Ser Gly Leu Ile  Gly Pro Leu Leu Val  Cys His Thr
        995                 1000                1005

Asn Thr  Leu Asn Pro Ala His  Gly Arg Gln Val Thr  Val Gln Glu
    1010                1015                1020

Phe Ala  Leu Phe Phe Thr Ile  Phe Asp Glu Thr Lys  Ser Trp Tyr
    1025                1030                1035

Phe Thr  Glu Asn Met Glu Arg  Asn Cys Arg Ala Pro  Cys Asn Ile
    1040                1045                1050

Gln Met  Glu Asp Pro Thr Phe  Lys Glu Asn Tyr Arg  Phe His Ala
    1055                1060                1065

Ile Asn  Gly Tyr Ile Met Asp  Thr Leu Pro Gly Leu  Val Met Ala
    1070                1075                1080

Gln Asp  Gln Arg Ile Arg Trp  Tyr Leu Leu Ser Met  Gly Ser Asn
    1085                1090                1095

Glu Asn  Ile His Ser Ile His  Phe Ser Gly His Val  Phe Thr Val
    1100                1105                1110

Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
    1115                1120                1125

Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    1130                1135                1140

Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    1145                1150                1155

Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    1160                1165                1170

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
    1175                1180                1185

Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
    1190                1195                1200

Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
    1205                1210                1215
```

```
Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
    1220                1225                1230

Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    1235                1240                1245

Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
    1250                1255                1260

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    1265                1270                1275

Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
    1280                1285                1290

Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1295                1300                1305

Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
    1310                1315                1320

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    1325                1330                1335

Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
    1340                1345                1350

Ser Lys  Ala Arg Leu His Leu  Gln Gly Arg Ser Asn  Ala Trp Arg
    1355                1360                1365

Pro Gln  Val Asn Asn Pro Lys  Glu Trp Leu Gln Val  Asp Phe Gln
    1370                1375                1380

Lys Thr  Met Lys Val Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser
    1385                1390                1395

Leu Leu  Thr Ser Met Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser
    1400                1405                1410

Gln Asp  Gly His Gln Trp Thr  Leu Phe Phe Gln Asn  Gly Lys Val
    1415                1420                1425

Lys Val  Phe Gln Gly Asn Gln  Asp Ser Phe Thr Pro  Val Val Asn
    1430                1435                1440

Ser Leu  Asp Pro Pro Leu Leu  Thr Arg Tyr Leu Arg  Ile His Pro
    1445                1450                1455

Gln Ser  Trp Val His Gln Ile  Ala Leu Arg Met Glu  Val Leu Gly
    1460                1465                1470
```

Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu
1475 1480 1485

Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala
1490 1495 1500

Thr Gly His His His His His His
1505 1510

<210> 648
<211> 4536
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

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agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

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<211> 1511
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val

65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
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Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp

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Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
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Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
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Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
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Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
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Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
740 745 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
755 760 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
770 775 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785 790 795 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
805 810 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
820 825 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
835 840 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
850 855 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865 870 875 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile

885 890 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Gly Ala Pro Gly
900 905 910

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
915 920 925

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
930 935 940

Glu Thr Pro Ala Ser Ser Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
945 950 955 960

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp
965 970 975

Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu
980 985 990

Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val Cys His Thr
995 1000 1005

Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr Val Gln Glu
1010 1015 1020

Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys Ser Trp Tyr
1025 1030 1035

Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro Cys Asn Ile
1040 1045 1050

Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg Phe His Ala
1055 1060 1065

Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu Val Met Ala
1070 1075 1080

Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met Gly Ser Asn
1085 1090 1095

Glu Asn Ile His Ser Ile His Phe Ser Gly His Val Phe Thr Val
1100 1105 1110

Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn Leu Tyr Pro
1115 1120 1125

Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys Ala Gly Ile
1130 1135 1140

Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His Ala Gly Met

1145 1150 1155

Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln Thr Pro Leu
1160 1165 1170

Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile Thr Ala Ser
1175 1180 1185

Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg Leu His Tyr
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Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro Phe Ser Trp
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Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His Gly Ile Lys
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Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr Ile Ser Gln
1235 1240 1245

Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp Gln Thr Tyr
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Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe Gly Asn Val
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Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro Pro Ile Ile
1280 1285 1290

Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser Ile Arg Ser
1295 1300 1305

Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn Ser Cys Ser
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Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp Ala Gln Ile
1325 1330 1335

Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp Ser Pro
1340 1345 1350

Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg
1355 1360 1365

Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln
1370 1375 1380

Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser
1385 1390 1395

Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser

1400 1405 1410

Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val
1415 1420 1425

Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn
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Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro
1445 1450 1455

Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly
1460 1465 1470

Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu
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Thr Gly His His His His His His
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Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1               5                  10                  15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270
```

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Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540
```

```
Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815
```

```
Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Gly Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
    930                 935                 940

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
945                 950                 955                 960

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Ala Ser Ser Asp
                965                 970                 975

Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu
            980                 985                 990

Lys Asp Val His Ser Gly Leu Ile  Gly Pro Leu Leu Val  Cys His Thr
        995                 1000                1005

Asn Thr  Leu Asn Pro Ala His  Gly Arg Gln Val Thr  Val Gln Glu
    1010                1015                1020

Phe Ala  Leu Phe Phe Thr Ile  Phe Asp Glu Thr Lys  Ser Trp Tyr
    1025                1030                1035

Phe Thr  Glu Asn Met Glu Arg  Asn Cys Arg Ala Pro  Cys Asn Ile
    1040                1045                1050

Gln Met  Glu Asp Pro Thr Phe  Lys Glu Asn Tyr Arg  Phe His Ala
    1055                1060                1065

Ile Asn  Gly Tyr Ile Met Asp  Thr Leu Pro Gly Leu  Val Met Ala
    1070                1075                1080
```

```
Gln Asp  Gln Arg Ile Arg Trp  Tyr Leu Leu Ser Met  Gly Ser Asn
    1085                 1090                 1095

Glu Asn  Ile His Ser Ile His  Phe Ser Gly His Val  Phe Thr Val
    1100                 1105                 1110

Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
    1115                 1120                 1125

Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    1130                 1135                 1140

Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    1145                 1150                 1155

Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    1160                 1165                 1170

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
    1175                 1180                 1185

Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
    1190                 1195                 1200

Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
    1205                 1210                 1215

Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
    1220                 1225                 1230

Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    1235                 1240                 1245

Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
    1250                 1255                 1260

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    1265                 1270                 1275

Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
    1280                 1285                 1290

Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1295                 1300                 1305

Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
    1310                 1315                 1320

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    1325                 1330                 1335
```

Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp Ser Pro
1340 1345 1350

Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg
1355 1360 1365

Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln
1370 1375 1380

Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser
1385 1390 1395

Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser
1400 1405 1410

Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val
1415 1420 1425

Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn
1430 1435 1440

Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro
1445 1450 1455

Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly
1460 1465 1470

Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu
1475 1480 1485

Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala
1490 1495 1500

Thr Gly His His His His His His
1505 1510

<210> 652
<211> 4536
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 652
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ctcggtgagc tgcctgtgga cgcaagattt cctcctagag tgccaaaatc ttttccattc 120

aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

agagatctag cttcaggact cattggccct ctcctcatct gctacaaaga atctgtagat 1680

caaagaggaa accagataat gtcagacaag aggaatgtca tcctgttttc tgtatttgat 1740

gagaaccgaa gctggtacct cacagagaat atacaacgct ttctccccaa tccagctgga 1800

gtgcagcttg aggatccaga gttccaagcc tccaacatca tgcacagcat caatggctat 1860

gtttttgata gtttgcagtt gtcagtttgt ttgcatgagg tggcatactg gtacattcta 1920

agcattggag cacagactga cttcctttct gtcttcttct ctggatatac cttcaaacac 1980

aaaatggtct atgaagacac actcacccta ttcccattct caggagaaac tgtcttcatg 2040

tcgatggaaa acccaggtct atggattctg gggtgccaca actcagactt tcggaacaga 2100

ggcatgaccg ccttactgaa ggtttctagt tgtgacaaga acactggtga ttattacgag 2160

gacagttatg aagatatttc agcatacttg ctgagtaaaa acaatgccat tgaaccaaga 2220

agcttctctc aaaacccacc agtcttgaaa cgccatcaac gggaaataac tcgtactact 2280

cttcagtcag atcaagagga aatcgattat gatgatacca tatcagttga aatgaagaag 2340

gaagattttg acatttatga tgaggatgaa aatcagagcc cccgcagctt tcaaaagaaa 2400

acacgacact attttattgc tgcagtggag aggctctggg attatgggat gagtagctcc 2460

ccacatgttc taagaaacag ggctcagagt ggcagtgtcc ctcagttcaa gaaagttgtt 2520

ttccaggaat ttactgatgg ctcctttact cagcccttat accgtggaga actaaatgaa 2580

catttgggac tcctggggcc atatataaga gcagaagttg aagataatat catggtaact 2640

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aaagagaatt atcgcttcca tgcaatcaat ggctacataa tggatacact acctggctta 3240

gtaatggctc aggatcaaag gattcgatgg tatctgctca gcatgggcag caatgaaaac 3300

atccattcta ttcatttcag tggacatgtg ttcactgtac gaaaaaaaga ggagtataaa 3360

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ctttttctgg tgtacagcaa taagtgtcag actcccctgg gaatggcttc tggacacatt 3540

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cagacttatc gaggaaattc cactggaacc ttaatggtct tctttggcaa tgtggattca 3840

tctgggataa aacacaatat ttttaaccct ccaattattg ctcgatacat ccgtttgcac 3900

ccaactcatt atagcattcg cagcactctt cgcatggagt tgatgggctg tgatttaaat 3960

agttgcagca tgccattggg aatggagagt aaagcaatat cagatgcaca gattactgct 4020

tcatcctact ttaccaatat gtttgccacc tggtctcctt caaaagctcg acttcacctc 4080

caagggagga gtaatgcctg gagacctcag gtgaataatc caaaagagtg gctgcaagtg 4140

gacttccaga agacaatgaa agtcacagga gtaactactc agggagtaaa atctctgctt 4200

accagcatgt atgtgaagga gttcctcatc tccagcagtc aagatggcca tcagtggact 4260

ctctttttc agaatggcaa agtaaaggtt tttcagggaa atcaagactc cttcacacct 4320

gtggtgaact ctctagaccc accgttactg actcgctacc ttcgaattca cccccagagt 4380

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<210> 653
<211> 1511
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 653

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

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His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480
```

```
Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750
```

```
Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Gly Ala Pro Gly Ser Pro Ala Gly Ser
                965                 970                 975

Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
            980                 985                 990

Gly Pro Gly Ser Glu Pro Ala Thr  Ser Gly Ser Glu Thr  Pro Ala Ser
        995                 1000                1005

Ser Thr  Leu Asn Pro Ala His  Gly Arg Gln Val Thr  Val Gln Glu
    1010                1015                1020
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Phe Ala  Leu Phe Phe Thr Ile  Phe Asp Glu Thr Lys  Ser Trp Tyr
    1025                1030                1035

Phe Thr  Glu Asn Met Glu Arg  Asn Cys Arg Ala Pro  Cys Asn Ile
    1040                1045                1050

Gln Met  Glu Asp Pro Thr Phe  Lys Glu Asn Tyr Arg  Phe His Ala
    1055                1060                1065

Ile Asn  Gly Tyr Ile Met Asp  Thr Leu Pro Gly Leu  Val Met Ala
    1070                1075                1080

Gln Asp  Gln Arg Ile Arg Trp  Tyr Leu Leu Ser Met  Gly Ser Asn
    1085                1090                1095

Glu Asn  Ile His Ser Ile His  Phe Ser Gly His Val  Phe Thr Val
    1100                1105                1110

Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
    1115                1120                1125

Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    1130                1135                1140

Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    1145                1150                1155

Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    1160                1165                1170

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
    1175                1180                1185

Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
    1190                1195                1200

Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
    1205                1210                1215

Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
    1220                1225                1230

Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    1235                1240                1245

Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
    1250                1255                1260

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    1265                1270                1275
```

```
Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
    1280                 1285                 1290

Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1295                 1300                 1305

Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
    1310                 1315                 1320

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    1325                 1330                 1335

Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
    1340                 1345                 1350

Ser Lys  Ala Arg Leu His Leu  Gln Gly Arg Ser Asn  Ala Trp Arg
    1355                 1360                 1365

Pro Gln  Val Asn Asn Pro Lys  Glu Trp Leu Gln Val  Asp Phe Gln
    1370                 1375                 1380

Lys Thr  Met Lys Val Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser
    1385                 1390                 1395

Leu Leu  Thr Ser Met Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser
    1400                 1405                 1410

Gln Asp  Gly His Gln Trp Thr  Leu Phe Phe Gln Asn  Gly Lys Val
    1415                 1420                 1425

Lys Val  Phe Gln Gly Asn Gln  Asp Ser Phe Thr Pro  Val Val Asn
    1430                 1435                 1440

Ser Leu  Asp Pro Pro Leu Leu  Thr Arg Tyr Leu Arg  Ile His Pro
    1445                 1450                 1455

Gln Ser  Trp Val His Gln Ile  Ala Leu Arg Met Glu  Val Leu Gly
    1460                 1465                 1470

Cys Glu  Ala Gln Asp Leu Tyr  Gly Ala Gly Ser Pro  Gly Ala Glu
    1475                 1480                 1485

Thr Ala  Glu Gln Lys Leu Ile  Ser Glu Glu Asp Leu  Ser Pro Ala
    1490                 1495                 1500

Thr Gly  His His His His His  His
    1505                 1510
```

<210> 654
<211> 4536
<212> DNA

<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

agagatctag cttcaggact cattggccct ctcctcatct gctacaaaga atctgtagat 1680

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gaagattttg acatttatga tgaggatgaa aatcagagcc cccgcagctt tcaaaagaaa 2400

acacgacact attttattgc tgcagtggag aggctctggg attatgggat gagtagctcc 2460

ccacatgttc taagaaacag ggctcagagt ggcagtgtcc ctcagttcaa gaaagttgtt 2520

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tgggcttatt tctctgatgt tgacctggaa aaagatgtgc actcaggcct gattggaccc 2880

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gtaatggctc aggatcaaag gattcgatgg tatctgctca gcatgggcag caatgaaaac 3300

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cagacttatc gaggaaattc cactggaacc ttaatggtct tctttggcaa tgtggattca 3840

tctgggataa aacacaatat ttttaaccct ccaattattg ctcgatacat ccgtttgcac 3900

ccaactcatt atagcattcg cagcactctt cgcatggagt tgatgggctg tgatttaaat 3960

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tcatcctact ttaccaatat gtttgccacc tggtctcctt caaaagctcg acttcacctc 4080

caagggagga gtaatgcctg gagacctcag gtgaataatc caaaagagtg gctgcaagtg 4140

gacttccaga agacaatgaa agtcacagga gtaactactc agggagtaaa atctctgctt 4200

accagcatgt atgtgaagga gttcctcatc tccagcagtc aagatggcca tcagtggact 4260

ctctttttc agaatggcaa agtaaaggtt tttcagggaa atcaagactc cttcacacct 4320

gtggtgaact ctctagaccc accgttactg actcgctacc ttcgaattca cccccagagt 4380

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tcacctgcaa ccggtcatca tcaccatcac cattga 4536

<210> 655
<211> 1511
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 655

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

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Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415
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Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685
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Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960
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Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Gly Ala Pro  Gly Ser Pro
        995                 1000                1005

Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Thr Ser  Glu Ser Ala
    1010                1015                1020

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1025                1030                1035

Glu Thr  Pro Ala Ser Ser Arg  Asn Cys Arg Ala Pro  Cys Asn Ile
    1040                1045                1050

Gln Met  Glu Asp Pro Thr Phe  Lys Glu Asn Tyr Arg  Phe His Ala
    1055                1060                1065

Ile Asn  Gly Tyr Ile Met Asp  Thr Leu Pro Gly Leu  Val Met Ala
    1070                1075                1080

Gln Asp  Gln Arg Ile Arg Trp  Tyr Leu Leu Ser Met  Gly Ser Asn
    1085                1090                1095

Glu Asn  Ile His Ser Ile His  Phe Ser Gly His Val  Phe Thr Val
    1100                1105                1110

Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
    1115                1120                1125

Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    1130                1135                1140

Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    1145                1150                1155

Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    1160                1165                1170

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
    1175                1180                1185

Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
    1190                1195                1200

Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
    1205                1210                1215
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```
Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
    1220                 1225                 1230

Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    1235                 1240                 1245

Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
    1250                 1255                 1260

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    1265                 1270                 1275

Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
    1280                 1285                 1290

Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1295                 1300                 1305

Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
    1310                 1315                 1320

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    1325                 1330                 1335

Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
    1340                 1345                 1350

Ser Lys  Ala Arg Leu His Leu  Gln Gly Arg Ser Asn  Ala Trp Arg
    1355                 1360                 1365

Pro Gln  Val Asn Asn Pro Lys  Glu Trp Leu Gln Val  Asp Phe Gln
    1370                 1375                 1380

Lys Thr  Met Lys Val Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser
    1385                 1390                 1395

Leu Leu  Thr Ser Met Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser
    1400                 1405                 1410

Gln Asp  Gly His Gln Trp Thr  Leu Phe Phe Gln Asn  Gly Lys Val
    1415                 1420                 1425

Lys Val  Phe Gln Gly Asn Gln  Asp Ser Phe Thr Pro  Val Val Asn
    1430                 1435                 1440

Ser Leu  Asp Pro Pro Leu Leu  Thr Arg Tyr Leu Arg  Ile His Pro
    1445                 1450                 1455

Gln Ser  Trp Val His Gln Ile  Ala Leu Arg Met Glu  Val Leu Gly
    1460                 1465                 1470
```

Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu
1475 1480 1485

Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala
1490 1495 1500

Thr Gly His His His His His His
1505 1510

<210> 656
<211> 4536
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

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catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

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gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

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aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

agagatctag cttcaggact cattggccct ctcctcatct gctacaaaga atctgtagat 1680

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gacagttatg aagatatttc agcatacttg ctgagtaaaa acaatgccat tgaaccaaga 2220

agcttctctc aaaacccacc agtcttgaaa cgccatcaac gggaaataac tcgtactact 2280

cttcagtcag atcaagagga aatcgattat gatgatacca tatcagttga aatgaagaag 2340

gaagattttg acatttatga tgaggatgaa aatcagagcc cccgcagctt tcaaaagaaa 2400

acacgacact attttattgc tgcagtggag aggctctggg attatgggat gagtagctcc 2460

ccacatgttc taagaaacag ggctcagagt ggcagtgtcc ctcagttcaa gaaagttgtt 2520

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gatcagaggc aaggagcaga acctagaaaa aactttgtca agcctaatga aaccaaaact 2760

tacttttgga aagtgcaaca tcatatggca cccactaaag atgagtttga ctgcaaagcc 2820

tgggcttatt tctctgatgt tgacctggaa aaagatgtgc actcaggcct gattggaccc 2880

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tcatcctact ttaccaatat gtttgccacc tggtctcctt caaaagctcg acttcacctc 4080

caagggagga gtaatgcctg gagacctcag gtgaataatc caaaagagtg gctgcaagtg 4140

gacttccaga agacaatgaa agtcacagga gtaactactc agggagtaaa atctctgctt 4200

accagcatgt atgtgaagga gttcctcatc tccagcagtc aagatggcca tcagtggact 4260

ctctttttc agaatggcaa agtaaaggtt tttcagggaa atcaagactc cttcacacct 4320

gtggtgaact ctctagaccc accgttactg actcgctacc ttcgaattca cccccagagt 4380

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tcacctgcaa ccggtcatca tcaccatcac cattga 4536

<210> 657
<211> 1511
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 657

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val

```
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
```

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            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
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610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
740 745 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
755 760 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
770 775 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785 790 795 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
805 810 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
820 825 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
835 840 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
850 855 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865 870 875 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile

885 890 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
900 905 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
915 920 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
930 935 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945 950 955 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
965 970 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
980 985 990

Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Gly Ala
995 1000 1005

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr
1010 1015 1020

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1025 1030 1035

Thr Ser Gly Ser Glu Thr Pro Ala Ser Ser Ala Pro Cys Asn Ile
1040 1045 1050

Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg Phe His Ala
1055 1060 1065

Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu Val Met Ala
1070 1075 1080

Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met Gly Ser Asn
1085 1090 1095

Glu Asn Ile His Ser Ile His Phe Ser Gly His Val Phe Thr Val
1100 1105 1110

Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn Leu Tyr Pro
1115 1120 1125

Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys Ala Gly Ile
1130 1135 1140

Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His Ala Gly Met

1145 1150 1155

Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln Thr Pro Leu
1160 1165 1170

Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile Thr Ala Ser
1175 1180 1185

Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg Leu His Tyr
1190 1195 1200

Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro Phe Ser Trp
1205 1210 1215

Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His Gly Ile Lys
1220 1225 1230

Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr Ile Ser Gln
1235 1240 1245

Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp Gln Thr Tyr
1250 1255 1260

Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe Gly Asn Val
1265 1270 1275

Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro Pro Ile Ile
1280 1285 1290

Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser Ile Arg Ser
1295 1300 1305

Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn Ser Cys Ser
1310 1315 1320

Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp Ala Gln Ile
1325 1330 1335

Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp Ser Pro
1340 1345 1350

Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg
1355 1360 1365

Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln
1370 1375 1380

Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser
1385 1390 1395

Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser

1400 1405 1410

Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val
1415 1420 1425

Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn
1430 1435 1440

Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro
1445 1450 1455

Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly
1460 1465 1470

Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu
1475 1480 1485

Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala
1490 1495 1500

Thr Gly His His His His His His
1505 1510

<210> 658
<211> 4536
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

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gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

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accagcatgt atgtgaagga gttcctcatc tccagcagtc aagatggcca tcagtggact 4260

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<210> 659
<211> 1511
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 659

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270
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Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540
```

```
Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815
```

```
Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Gly Ala Pro Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr
    1010                1015                1020

Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    1025                1030                1035

Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr Pro Ala  Ser Ser Ile
    1040                1045                1050

Gln Met  Glu Asp Pro Thr Phe  Lys Glu Asn Tyr Arg  Phe His Ala
    1055                1060                1065

Ile Asn  Gly Tyr Ile Met Asp  Thr Leu Pro Gly Leu  Val Met Ala
    1070                1075                1080
```

```
Gln Asp  Gln Arg Ile Arg Trp  Tyr Leu Leu Ser Met  Gly Ser Asn
    1085                 1090                 1095

Glu Asn  Ile His Ser Ile His  Phe Ser Gly His Val  Phe Thr Val
    1100                 1105                 1110

Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
    1115                 1120                 1125

Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    1130                 1135                 1140

Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    1145                 1150                 1155

Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    1160                 1165                 1170

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
    1175                 1180                 1185

Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
    1190                 1195                 1200

Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
    1205                 1210                 1215

Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
    1220                 1225                 1230

Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    1235                 1240                 1245

Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
    1250                 1255                 1260

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    1265                 1270                 1275

Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
    1280                 1285                 1290

Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1295                 1300                 1305

Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
    1310                 1315                 1320

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    1325                 1330                 1335
```

Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp Ser Pro
1340 1345 1350

Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg
1355 1360 1365

Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln
1370 1375 1380

Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser
1385 1390 1395

Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser
1400 1405 1410

Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val
1415 1420 1425

Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn
1430 1435 1440

Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro
1445 1450 1455

Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly
1460 1465 1470

Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu
1475 1480 1485

Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala
1490 1495 1500

Thr Gly His His His His His His
1505 1510

<210> 660
<211> 4536
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 660
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ctcggtgagc tgcctgtgga cgcaagattt cctcctagag tgccaaaatc ttttccattc 120

aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

agagatctag cttcaggact cattggccct ctcctcatct gctacaaaga atctgtagat 1680

caaagaggaa accagataat gtcagacaag aggaatgtca tcctgttttc tgtatttgat 1740

gagaaccgaa gctggtacct cacagagaat atacaacgct ttctccccaa tccagctgga 1800

gtgcagcttg aggatccaga gttccaagcc tccaacatca tgcacagcat caatggctat 1860

gtttttgata gtttgcagtt gtcagtttgt ttgcatgagg tggcatactg gtacattcta 1920

agcattggag cacagactga cttcctttct gtcttcttct ctggatatac cttcaaacac 1980

aaaatggtct atgaagacac actcacccta ttcccattct caggagaaac tgtcttcatg 2040

tcgatggaaa acccaggtct atggattctg gggtgccaca actcagactt tcggaacaga 2100

ggcatgaccg ccttactgaa ggtttctagt tgtgacaaga acactggtga ttattacgag 2160

gacagttatg aagatatttc agcatacttg ctgagtaaaa acaatgccat tgaaccaaga 2220

agcttctctc aaaacccacc agtcttgaaa cgccatcaac gggaaataac tcgtactact 2280

cttcagtcag atcaagagga aatcgattat gatgatacca tatcagttga aatgaagaag 2340

gaagattttg acatttatga tgaggatgaa aatcagagcc cccgcagctt tcaaaagaaa 2400

acacgacact attttattgc tgcagtggag aggctctggg attatgggat gagtagctcc 2460

ccacatgttc taagaaacag ggctcagagt ggcagtgtcc ctcagttcaa gaaagttgtt 2520

ttccaggaat ttactgatgg ctcctttact cagcccttat accgtggaga actaaatgaa 2580

catttgggac tcctggggcc atatataaga gcagaagttg aagataatat catggtaact 2640

ttcagaaatc aggcctctcg tccctattcc ttctattcta gccttatttc ttatgaggaa 2700

gatcagaggc aaggagcaga acctagaaaa aactttgtca agcctaatga aaccaaaact 2760

tacttttgga aagtgcaaca tcatatggca cccactaaag atgagtttga ctgcaaagcc 2820

tgggcttatt tctctgatgt tgacctggaa aaagatgtgc actcaggcct gattggaccc 2880

cttctggtct gccacactaa cacactgaac cctgctcatg ggagacaagt gacagtacag 2940

gaatttgctc tgtttttcac catctttgat gagaccaaaa gctggtactt cactgaaaat 3000

atggaaagaa actgcagggc tccctgcaat ggcgcgccag gttctcctgc tggctccccc 3060

acctcaacag aagaggggac aagcgaaagc gctacgcctg agagtggccc tggctctgag 3120

ccagccacct ccggctctga aacccctgcc tcgagcatcc agatggaaga tcccactttt 3180

aaagagaatt atcgcttcca tgcaatcaat ggctacataa tggatacact acctggctta 3240

gtaatggctc aggatcaaag gattcgatgg tatctgctca gcatgggcag caatgaaaac 3300

atccattcta ttcatttcag tggacatgtg ttcactgtac gaaaaaaaga ggagtataaa 3360

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cagacttatc gaggaaattc cactggaacc ttaatggtct tctttggcaa tgtggattca 3840

tctgggataa aacacaatat ttttaaccct ccaattattg ctcgatacat ccgtttgcac 3900

ccaactcatt atagcattcg cagcactctt cgcatggagt tgatgggctg tgatttaaat 3960

agttgcagca tgccattggg aatggagagt aaagcaatat cagatgcaca gattactgct 4020

tcatcctact ttaccaatat gtttgccacc tggtctcctt caaaagctcg acttcacctc 4080

caagggagga gtaatgcctg gagacctcag gtgaataatc caaaagagtg gctgcaagtg 4140

gacttccaga agacaatgaa agtcacagga gtaactactc agggagtaaa atctctgctt 4200

accagcatgt atgtgaagga gttcctcatc tccagcagtc aagatggcca tcagtggact 4260

ctctttttc agaatggcaa agtaaaggtt tttcagggaa atcaagactc cttcacacct 4320

gtggtgaact ctctagaccc accgttactg actcgctacc ttcgaattca cccccagagt 4380

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tcacctgcaa ccggtcatca tcaccatcac cattga 4536

<210> 661
<211> 1511
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 661

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

```
His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480
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Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750
```

```
Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020
```

```
Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1025                1030                1035

Val Met  Ala Gln Asp Gly Ala  Pro Gly Ser Pro Ala  Gly Ser Pro
    1040                1045                1050

Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1055                1060                1065

Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Ala
    1070                1075                1080

Ser Ser  Gln Arg Ile Arg Trp  Tyr Leu Leu Ser Met  Gly Ser Asn
    1085                1090                1095

Glu Asn  Ile His Ser Ile His  Phe Ser Gly His Val  Phe Thr Val
    1100                1105                1110

Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
    1115                1120                1125

Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    1130                1135                1140

Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    1145                1150                1155

Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    1160                1165                1170

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
    1175                1180                1185

Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
    1190                1195                1200

Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
    1205                1210                1215

Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
    1220                1225                1230

Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    1235                1240                1245

Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
    1250                1255                1260

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    1265                1270                1275
```

```
Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
    1280                1285                1290

Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1295                1300                1305

Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
    1310                1315                1320

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    1325                1330                1335

Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
    1340                1345                1350

Ser Lys  Ala Arg Leu His Leu  Gln Gly Arg Ser Asn  Ala Trp Arg
    1355                1360                1365

Pro Gln  Val Asn Asn Pro Lys  Glu Trp Leu Gln Val  Asp Phe Gln
    1370                1375                1380

Lys Thr  Met Lys Val Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser
    1385                1390                1395

Leu Leu  Thr Ser Met Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser
    1400                1405                1410

Gln Asp  Gly His Gln Trp Thr  Leu Phe Phe Gln Asn  Gly Lys Val
    1415                1420                1425

Lys Val  Phe Gln Gly Asn Gln  Asp Ser Phe Thr Pro  Val Val Asn
    1430                1435                1440

Ser Leu  Asp Pro Pro Leu Leu  Thr Arg Tyr Leu Arg  Ile His Pro
    1445                1450                1455

Gln Ser  Trp Val His Gln Ile  Ala Leu Arg Met Glu  Val Leu Gly
    1460                1465                1470

Cys Glu  Ala Gln Asp Leu Tyr  Gly Ala Gly Ser Pro  Gly Ala Glu
    1475                1480                1485

Thr Ala  Glu Gln Lys Leu Ile  Ser Glu Glu Asp Leu  Ser Pro Ala
    1490                1495                1500

Thr Gly  His His His His His  His
    1505                1510
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<210> 662
<211> 4536
<212> DNA

<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 662
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<213> Artificial Sequence

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

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Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415
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Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
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Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
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Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685
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Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
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Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
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Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
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Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
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Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
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Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
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Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1055                1060                1065

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1070                1075                1080

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1085                1090                1095

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
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Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Gly Ala Pro
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Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser
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Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
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Ser Gly  Ser Glu Thr Pro Ala  Ser Ser Lys Cys Gln  Thr Pro Leu
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Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
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Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
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Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
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Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
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Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    1235                 1240                 1245

Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
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Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
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Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
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Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
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Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
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Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
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Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
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Ser Lys  Ala Arg Leu His Leu  Gln Gly Arg Ser Asn  Ala Trp Arg
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Pro Gln  Val Asn Asn Pro Lys  Glu Trp Leu Gln Val  Asp Phe Gln
    1370                 1375                 1380

Lys Thr  Met Lys Val Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser
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Leu Leu  Thr Ser Met Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser
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Gln Asp  Gly His Gln Trp Thr  Leu Phe Phe Gln Asn  Gly Lys Val
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Lys Val  Phe Gln Gly Asn Gln  Asp Ser Phe Thr Pro  Val Val Asn
    1430                 1435                 1440

Ser Leu  Asp Pro Pro Leu Leu  Thr Arg Tyr Leu Arg  Ile His Pro
    1445                 1450                 1455

Gln Ser  Trp Val His Gln Ile  Ala Leu Arg Met Glu  Val Leu Gly
    1460                 1465                 1470
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Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu
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Thr Gly His His His His His His
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ccaactcatt atagcattcg cagcactctt cgcatggagt tgatgggctg tgatttaaat 3960

agttgcagca tgccattggg aatggagagt aaagcaatat cagatgcaca gattactgct 4020

tcatcctact ttaccaatat gtttgccacc tggtctcctt caaaagctcg acttcacctc 4080

caagggagga gtaatgcctg gagacctcag gtgaataatc caaaagagtg gctgcaagtg 4140

gacttccaga agacaatgaa agtcacagga gtaactactc agggagtaaa atctctgctt 4200

accagcatgt atgtgaagga gttcctcatc tccagcagtc aagatggcca tcagtggact 4260

ctctttttc agaatggcaa agtaaaggtt tttcagggaa atcaagactc cttcacacct 4320

gtggtgaact ctctagaccc accgttactg actcgctacc ttcgaattca cccccagagt 4380

tgggtgcacc agattgccct gaggatggag gttctgggct gcgaggcaca ggacctctac 4440

ggcgccggat cacctggggc cgaaacggcc gaacaaaaac tcatctcaga agaggatctg 4500

tcacctgcaa ccggtcatca tcaccatcac cattga 4536

<210> 665
<211> 1511
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 665
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val

65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp

```
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
```

610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
740 745 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
755 760 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
770 775 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785 790 795 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
805 810 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
820 825 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
835 840 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
850 855 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865 870 875 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile

885 890 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
900 905 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
915 920 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
930 935 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945 950 955 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
965 970 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
980 985 990

Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
995 1000 1005

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1010 1015 1020

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1025 1030 1035

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1040 1045 1050

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1055 1060 1065

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1070 1075 1080

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1085 1090 1095

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1100 1105 1110

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1115 1120 1125

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1130 1135 1140

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg

```
    1145                1150                1155

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1160                1165                1170

Phe Gly  Ala Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    1175                1180                1185

Glu Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1190                1195                1200

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Ala Ser  Ser Ser Trp
    1205                1210                1215

Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
    1220                1225                1230

Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    1235                1240                1245

Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
    1250                1255                1260

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    1265                1270                1275

Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
    1280                1285                1290

Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1295                1300                1305

Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
    1310                1315                1320

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    1325                1330                1335

Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
    1340                1345                1350

Ser Lys  Ala Arg Leu His Leu  Gln Gly Arg Ser Asn  Ala Trp Arg
    1355                1360                1365

Pro Gln  Val Asn Asn Pro Lys  Glu Trp Leu Gln Val  Asp Phe Gln
    1370                1375                1380

Lys Thr  Met Lys Val Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser
    1385                1390                1395

Leu Leu  Thr Ser Met Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser
```

1400 1405 1410

Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val
1415 1420 1425

Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn
1430 1435 1440

Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro
1445 1450 1455

Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly
1460 1465 1470

Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu
1475 1480 1485

Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala
1490 1495 1500

Thr Gly His His His His His His
1505 1510

<210> 666
<211> 4536
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 666
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ctcggtgagc tgcctgtgga cgcaagattt cctcctagag tgccaaaatc ttttccattc 120

aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

agagatctag cttcaggact cattggccct ctcctcatct gctacaaaga atctgtagat 1680

caaagaggaa accagataat gtcagacaag aggaatgtca tcctgttttc tgtatttgat 1740

gagaaccgaa gctggtacct cacagagaat atacaacgct ttctccccaa tccagctgga 1800

gtgcagcttg aggatccaga gttccaagcc tccaacatca tgcacagcat caatggctat 1860

gtttttgata gtttgcagtt gtcagtttgt ttgcatgagg tggcatactg gtacattcta 1920

agcattggag cacagactga cttccttttct gtcttcttct ctggatatac cttcaaacac 1980

aaaatggtct atgaagacac actcacccta ttcccattct caggagaaac tgtcttcatg 2040

tcgatggaaa acccaggtct atggattctg gggtgccaca actcagactt tcggaacaga 2100

ggcatgaccg ccttactgaa ggtttctagt tgtgacaaga acactggtga ttattacgag 2160

gacagttatg aagatatttc agcatacttg ctgagtaaaa acaatgccat tgaaccaaga 2220

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cttcagtcag atcaagagga aatcgattat gatgatacca tatcagttga aatgaagaag 2340

gaagattttg acatttatga tgaggatgaa aatcagagcc cccgcagctt tcaaaagaaa 2400

acacgacact attttattgc tgcagtggag aggctctggg attatgggat gagtagctcc 2460

ccacatgttc taagaaacag ggctcagagt ggcagtgtcc ctcagttcaa gaaagttgtt 2520

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gaatttgctc tgtttttcac catctttgat gagaccaaaa gctggtactt cactgaaaat 3000

atggaaagaa actgcagggc tccctgcaat atccagatgg aagatcccac ttttaaagag 3060

aattatcgct tccatgcaat caatggctac ataatggata cactacctgg cttagtaatg 3120

gctcaggatc aaaggattcg atggtatctg ctcagcatgg gcagcaatga aaacatccat 3180

tctattcatt tcagtggaca tgtgttcact gtacgaaaaa aagaggagta taaaatggca 3240

ctgtacaatc tctatccagg tgtttttgag acagtggaaa tgttaccatc caaagctgga 3300

atttggcggg tggaatgcct tattggcgag catctacatg ctgggatgag cacacttttt 3360

ctggtgtaca gcaataagtg tcagactccc ctgggaatgg cttctggaca cattagagat 3420

tttcagatta cagcttcagg acaatatgga cagtgggccc caaagctggc cagacttcat 3480

tattccggat caatcaatgc ctggagcacc aaggagccct ttggcgcgcc aggttctcct 3540

gctggctccc ccacctcaac agaagagggg acaagcgaaa gcgctacgcc tgagagtggc 3600

cctggctctg agccagccac ctccggctct gaaacccctg cctcgagctc ttggatcaag 3660

gtggatctgt tggcaccaat gattattcac ggcatcaaga cccagggtgc ccgtcagaag 3720

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cagacttatc gaggaaattc cactggaacc ttaatggtct tctttggcaa tgtggattca 3840

tctgggataa aacacaatat ttttaaccct ccaattattg ctcgatacat ccgtttgcac 3900

ccaactcatt atagcattcg cagcactctt cgcatggagt tgatgggctg tgatttaaat 3960

agttgcagca tgccattggg aatggagagt aaagcaatat cagatgcaca gattactgct 4020

tcatcctact ttaccaatat gtttgccacc tggtctcctt caaaagctcg acttcacctc 4080

caagggagga gtaatgcctg gagacctcag gtgaataatc caaaagagtg gctgcaagtg 4140

gacttccaga agacaatgaa agtcacagga gtaactactc agggagtaaa atctctgctt 4200

accagcatgt atgtgaagga gttcctcatc tccagcagtc aagatggcca tcagtggact 4260

ctctttttc agaatggcaa agtaaaggtt tttcagggaa atcaagactc cttcacacct 4320

gtggtgaact ctctagaccc accgttactg actcgctacc ttcgaattca cccccagagt 4380

tgggtgcacc agattgccct gaggatggag gttctgggct gcgaggcaca ggacctctac 4440

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tcacctgcaa ccggtcatca tcaccatcac cattga 4536

<210> 667
<211> 1511
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 667

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Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1               5                  10                  15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270
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Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540
```

```
Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815
```

```
Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1025                1030                1035

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1040                1045                1050

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1055                1060                1065

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1070                1075                1080
```

```
Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1085                1090                1095

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1100                1105                1110

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1115                1120                1125

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1130                1135                1140

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1145                1150                1155

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1160                1165                1170

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1175                1180                1185

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1190                1195                1200

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Gly
    1205                1210                1215

Ala Pro  Gly Ser Pro Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly
    1220                1225                1230

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro
    1235                1240                1245

Ala Thr  Ser Gly Ser Glu Thr  Pro Ala Ser Ser Trp  Gln Thr Tyr
    1250                1255                1260

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    1265                1270                1275

Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
    1280                1285                1290

Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1295                1300                1305

Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
    1310                1315                1320

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    1325                1330                1335
```

Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp Ser Pro
1340 1345 1350

Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg
1355 1360 1365

Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln
1370 1375 1380

Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser
1385 1390 1395

Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser
1400 1405 1410

Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val
1415 1420 1425

Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn
1430 1435 1440

Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro
1445 1450 1455

Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly
1460 1465 1470

Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu
1475 1480 1485

Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala
1490 1495 1500

Thr Gly His His His His His His
1505 1510

<210> 668
<211> 4536
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 668
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ctcggtgagc tgcctgtgga cgcaagattt cctcctagag tgccaaaatc ttttccattc 120

aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

agagatctag cttcaggact cattggccct ctcctcatct gctacaaaga atctgtagat 1680

caaagaggaa accagataat gtcagacaag aggaatgtca tcctgttttc tgtatttgat 1740

gagaaccgaa gctggtacct cacagagaat atacaacgct ttctccccaa tccagctgga 1800

gtgcagcttg aggatccaga gttccaagcc tccaacatca tgcacagcat caatggctat 1860

gtttttgata gtttgcagtt gtcagtttgt ttgcatgagg tggcatactg gtacattcta 1920

agcattggag cacagactga cttcctttct gtcttcttct ctggatatac cttcaaacac 1980

aaaatggtct atgaagacac actcacccta ttcccattct caggagaaac tgtcttcatg 2040

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ggcatgaccg ccttactgaa ggtttctagt tgtgacaaga acactggtga ttattacgag 2160

gacagttatg aagatatttc agcatacttg ctgagtaaaa acaatgccat tgaaccaaga 2220

agcttctctc aaaacccacc agtcttgaaa cgccatcaac gggaaataac tcgtactact 2280

cttcagtcag atcaagagga aatcgattat gatgatacca tatcagttga aatgaagaag 2340

gaagattttg acatttatga tgaggatgaa aatcagagcc cccgcagctt tcaaaagaaa 2400

acacgacact attttattgc tgcagtggag aggctctggg attatgggat gagtagctcc 2460

ccacatgttc taagaaacag ggctcagagt ggcagtgtcc ctcagttcaa gaaagttgtt 2520

ttccaggaat ttactgatgg ctcctttact cagcccttat accgtggaga actaaatgaa 2580

catttgggac tcctggggcc atatataaga gcagaagttg aagataatat catggtaact 2640

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gatcagaggc aaggagcaga acctagaaaa aactttgtca agcctaatga aaccaaaact 2760

tacttttgga aagtgcaaca tcatatggca cccactaaag atgagtttga ctgcaaagcc 2820

tgggcttatt tctctgatgt tgacctggaa aaagatgtgc actcaggcct gattggaccc 2880

cttctggtct gccacactaa cacactgaac cctgctcatg ggagacaagt gacagtacag 2940

gaatttgctc tgtttttcac catctttgat gagaccaaaa gctggtactt cactgaaaat 3000

atggaaagaa actgcagggc tccctgcaat atccagatgg aagatcccac ttttaaagag 3060

aattatcgct tccatgcaat caatggctac ataatggata cactacctgg cttagtaatg 3120

gctcaggatc aaaggattcg atggtatctg ctcagcatgg gcagcaatga aaacatccat 3180

tctattcatt tcagtggaca tgtgttcact gtacgaaaaa aagaggagta taaaatggca 3240

ctgtacaatc tctatccagg tgtttttgag acagtggaaa tgttaccatc caaagctgga 3300

atttggcggg tggaatgcct tattggcgag catctacatg ctgggatgag cacacttttt 3360

ctggtgtaca gcaataagtg tcagactccc ctgggaatgg cttctggaca cattagagat 3420

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tattccggat caatcaatgc ctggagcacc aaggagccct tttcttggat caaggtggat 3540

ctgttggcac caatgattat tcacggcatc aagacccagg gtgcccgtca gaagttctcc 3600

agcctctaca tctctcagtt tatcatcatg tatagtcttg atgggaagaa gggcgcgcca 3660

ggttctcctg ctggctcccc cacctcaaca gaagagggga caagcgaaag cgctacgcct 3720

gagagtggcc ctggctctga gccagccacc tccggctctg aaacccctgc ctcgagctgg 3780

cagacttatc gaggaaattc cactggaacc ttaatggtct tctttggcaa tgtggattca 3840

tctgggataa aacacaatat ttttaaccct ccaattattg ctcgatacat ccgtttgcac 3900

ccaactcatt atagcattcg cagcactctt cgcatggagt tgatgggctg tgatttaaat 3960

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tcatcctact ttaccaatat gtttgccacc tggtctcctt caaaagctcg acttcacctc 4080

caagggagga gtaatgcctg gagacctcag gtgaataatc caaaagagtg gctgcaagtg 4140

gacttccaga agacaatgaa agtcacagga gtaactactc agggagtaaa atctctgctt 4200

accagcatgt atgtgaagga gttcctcatc tccagcagtc aagatggcca tcagtggact 4260

ctctttttc agaatggcaa agtaaaggtt tttcagggaa atcaagactc cttcacacct 4320

gtggtgaact ctctagaccc accgttactg actcgctacc ttcgaattca cccccagagt 4380

tgggtgcacc agattgccct gaggatggag gttctgggct gcgaggcaca ggacctctac 4440

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tcacctgcaa ccggtcatca tcaccatcac cattga 4536

<210> 669
<211> 1511
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 669

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

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His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480
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Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750
```

```
Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020
```

```
Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1025                1030                1035

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1040                1045                1050

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1055                1060                1065

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1070                1075                1080

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1085                1090                1095

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1100                1105                1110

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1115                1120                1125

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1130                1135                1140

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1145                1150                1155

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1160                1165                1170

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1175                1180                1185

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1190                1195                1200

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1205                1210                1215

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Ala Pro Gly  Ser Pro Ala
    1220                1225                1230

Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr Ser Glu  Ser Ala Thr
    1235                1240                1245

Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu
    1250                1255                1260

Thr Pro  Ala Ser Ser Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    1265                1270                1275
```

```
Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
    1280                1285                1290

Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1295                1300                1305

Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
    1310                1315                1320

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    1325                1330                1335

Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
    1340                1345                1350

Ser Lys  Ala Arg Leu His Leu  Gln Gly Arg Ser Asn  Ala Trp Arg
    1355                1360                1365

Pro Gln  Val Asn Asn Pro Lys  Glu Trp Leu Gln Val  Asp Phe Gln
    1370                1375                1380

Lys Thr  Met Lys Val Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser
    1385                1390                1395

Leu Leu  Thr Ser Met Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser
    1400                1405                1410

Gln Asp  Gly His Gln Trp Thr  Leu Phe Phe Gln Asn  Gly Lys Val
    1415                1420                1425

Lys Val  Phe Gln Gly Asn Gln  Asp Ser Phe Thr Pro  Val Val Asn
    1430                1435                1440

Ser Leu  Asp Pro Pro Leu Leu  Thr Arg Tyr Leu Arg  Ile His Pro
    1445                1450                1455

Gln Ser  Trp Val His Gln Ile  Ala Leu Arg Met Glu  Val Leu Gly
    1460                1465                1470

Cys Glu  Ala Gln Asp Leu Tyr  Gly Ala Gly Ser Pro  Gly Ala Glu
    1475                1480                1485

Thr Ala  Glu Gln Lys Leu Ile  Ser Glu Glu Asp Leu  Ser Pro Ala
    1490                1495                1500

Thr Gly  His His His His His  His
    1505                1510
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<210> 670
<211> 4536
<212> DNA

<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 670
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ctcggtgagc tgcctgtgga cgcaagattt cctcctagag tgccaaaatc ttttccattc 120

aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

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gtgcagcttg aggatccaga gttccaagcc tccaacatca tgcacagcat caatggctat 1860

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<210> 671
<211> 1511
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 671

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

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Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415
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Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685
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Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960
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Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1025                1030                1035

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1040                1045                1050

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1055                1060                1065

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1070                1075                1080

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1085                1090                1095

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1100                1105                1110

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1115                1120                1125

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1130                1135                1140

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1145                1150                1155

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1160                1165                1170

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1175                1180                1185

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1190                1195                1200

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1205                1210                1215
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Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1220                 1225                 1230

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1235                 1240                 1245

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1250                 1255                 1260

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Gly
    1265                 1270                 1275

Ala Pro  Gly Ser Pro Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly
    1280                 1285                 1290

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro
    1295                 1300                 1305

Ala Thr  Ser Gly Ser Glu Thr  Pro Ala Ser Ser Asn  Ser Cys Ser
    1310                 1315                 1320

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    1325                 1330                 1335

Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
    1340                 1345                 1350

Ser Lys  Ala Arg Leu His Leu  Gln Gly Arg Ser Asn  Ala Trp Arg
    1355                 1360                 1365

Pro Gln  Val Asn Asn Pro Lys  Glu Trp Leu Gln Val  Asp Phe Gln
    1370                 1375                 1380

Lys Thr  Met Lys Val Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser
    1385                 1390                 1395

Leu Leu  Thr Ser Met Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser
    1400                 1405                 1410

Gln Asp  Gly His Gln Trp Thr  Leu Phe Phe Gln Asn  Gly Lys Val
    1415                 1420                 1425

Lys Val  Phe Gln Gly Asn Gln  Asp Ser Phe Thr Pro  Val Val Asn
    1430                 1435                 1440

Ser Leu  Asp Pro Pro Leu Leu  Thr Arg Tyr Leu Arg  Ile His Pro
    1445                 1450                 1455

Gln Ser  Trp Val His Gln Ile  Ala Leu Arg Met Glu  Val Leu Gly
    1460                 1465                 1470
```

Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu
1475 1480 1485

Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala
1490 1495 1500

Thr Gly His His His His His His
1505 1510

<210> 672
<211> 4536
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

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gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

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gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

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<210> 673
<211> 1511
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 673
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val

65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp

340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser

610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
740 745 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
755 760 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
770 775 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785 790 795 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
805 810 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
820 825 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
835 840 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
850 855 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865 870 875 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile

885 890 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
900 905 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
915 920 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
930 935 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945 950 955 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
965 970 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
980 985 990

Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
995 1000 1005

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1010 1015 1020

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1025 1030 1035

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1040 1045 1050

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1055 1060 1065

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1070 1075 1080

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1085 1090 1095

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1100 1105 1110

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1115 1120 1125

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1130 1135 1140

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg

1145 1150 1155

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1160 1165 1170

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1175 1180 1185

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1190 1195 1200

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1205 1210 1215

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1220 1225 1230

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1235 1240 1245

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1250 1255 1260

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1265 1270 1275

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1280 1285 1290

Ala Gly Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
1295 1300 1305

Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
1310 1315 1320

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Ala Ser Ser Gln Ile
1325 1330 1335

Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp Ser Pro
1340 1345 1350

Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg
1355 1360 1365

Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln
1370 1375 1380

Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser
1385 1390 1395

Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser

1400 1405 1410

Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val
1415 1420 1425

Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn
1430 1435 1440

Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro
1445 1450 1455

Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly
1460 1465 1470

Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu
1475 1480 1485

Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala
1490 1495 1500

Thr Gly His His His His His His
1505 1510

<210> 674
<211> 4536
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 674
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ctcggtgagc tgcctgtgga cgcaagattt cctcctagag tgccaaaatc ttttccattc 120

aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

agagatctag cttcaggact cattggccct ctcctcatct gctacaaaga atctgtagat 1680

caaagaggaa accagataat gtcagacaag aggaatgtca tcctgttttc tgtatttgat 1740

gagaaccgaa gctggtacct cacagagaat atacaacgct ttctccccaa tccagctgga 1800

gtgcagcttg aggatccaga gttccaagcc tccaacatca tgcacagcat caatggctat 1860

gtttttgata gtttgcagtt gtcagtttgt ttgcatgagg tggcatactg gtacattcta 1920

agcattggag cacagactga cttccttttct gtcttcttct ctggatatac cttcaaacac 1980

aaaatggtct atgaagacac actcacccta ttcccattct caggagaaac tgtcttcatg 2040

tcgatggaaa acccaggtct atggattctg gggtgccaca actcagactt tcggaacaga 2100

ggcatgaccg ccttactgaa ggtttctagt tgtgacaaga acactggtga ttattacgag 2160

gacagttatg aagatatttc agcatacttg ctgagtaaaa acaatgccat tgaaccaaga 2220

agcttctctc aaaacccacc agtcttgaaa cgccatcaac gggaaataac tcgtactact 2280

cttcagtcag atcaagagga aatcgattat gatgatacca tatcagttga aatgaagaag 2340

gaagattttg acatttatga tgaggatgaa aatcagagcc cccgcagctt tcaaaagaaa 2400

acacgacact attttattgc tgcagtggag aggctctggg attatgggat gagtagctcc 2460

ccacatgttc taagaaacag ggctcagagt ggcagtgtcc ctcagttcaa gaaagttgtt 2520

ttccaggaat ttactgatgg ctcctttact cagcccttat accgtggaga actaaatgaa 2580

catttgggac tcctggggcc atatataaga gcagaagttg aagataatat catggtaact 2640

ttcagaaatc aggcctctcg tccctattcc ttctattcta gccttatttc ttatgaggaa 2700

gatcagaggc aaggagcaga acctagaaaa aactttgtca agcctaatga aaccaaaact 2760

tacttttgga aagtgcaaca tcatatggca cccactaaag atgagtttga ctgcaaagcc 2820

tgggcttatt tctctgatgt tgacctggaa aaagatgtgc actcaggcct gattggaccc 2880

cttctggtct gccacactaa cacactgaac cctgctcatg ggagacaagt gacagtacag 2940

gaatttgctc tgtttttcac catctttgat gagaccaaaa gctggtactt cactgaaaat 3000

atggaaagaa actgcagggc tccctgcaat atccagatgg aagatcccac ttttaaagag 3060

aattatcgct tccatgcaat caatggctac ataatggata cactacctgg cttagtaatg 3120

gctcaggatc aaaggattcg atggtatctg ctcagcatgg gcagcaatga aaacatccat 3180

tctattcatt tcagtggaca tgtgttcact gtacgaaaaa aagaggagta taaaatggca 3240

ctgtacaatc tctatccagg tgtttttgag acagtggaaa tgttaccatc caaagctgga 3300

atttggcggg tggaatgcct tattggcgag catctacatg ctgggatgag cacacttttt 3360

ctggtgtaca gcaataagtg tcagactccc ctgggaatgg cttctggaca cattagagat 3420

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tattccggat caatcaatgc ctggagcacc aaggagccct tttcttggat caaggtggat 3540

ctgttggcac caatgattat tcacggcatc aagacccagg gtgcccgtca gaagttctcc 3600

agcctctaca tctctcagtt tatcatcatg tatagtcttg atgggaagaa gtggcagact 3660

tatcgaggaa attccactgg aaccttaatg gtcttctttg gcaatgtgga ttcatctggg 3720

ataaaacaca atatttttaa ccctccaatt attgctcgat acatccgttt gcacccaact 3780

cattatagca ttcgcagcac tcttcgcatg gagttgatgg gctgtgattt aaatagttgc 3840

agcatgccat tgggaatgga gagtaaagca atatcagatg caggcgcgcc aggttctcct 3900

gctggctccc ccacctcaac agaagagggg acaagcgaaa gcgctacgcc tgagagtggc 3960

cctggctctg agccagccac ctccggctct gaaacccctg cctcgagcca gattactgct 4020

tcatcctact ttaccaatat gtttgccacc tggtctcctt caaaagctcg acttcacctc 4080

caagggagga gtaatgcctg gagacctcag gtgaataatc caaaagagtg gctgcaagtg 4140

gacttccaga agacaatgaa agtcacagga gtaactactc agggagtaaa atctctgctt 4200

accagcatgt atgtgaagga gttcctcatc tccagcagtc aagatggcca tcagtggact 4260

ctctttttc agaatggcaa agtaaaggtt tttcagggaa atcaagactc cttcacacct 4320

gtggtgaact ctctagaccc accgttactg actcgctacc ttcgaattca cccccagagt 4380

tgggtgcacc agattgccct gaggatggag gttctgggct gcgaggcaca ggacctctac 4440

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tcacctgcaa ccggtcatca tcaccatcac cattga 4536

<210> 675
<211> 1511
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 675

```
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1               5                  10                  15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270
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Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540
```

```
Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815
```

```
Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1025                1030                1035

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1040                1045                1050

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1055                1060                1065

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1070                1075                1080
```

```
Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1085                1090                1095

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1100                1105                1110

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1115                1120                1125

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1130                1135                1140

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1145                1150                1155

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1160                1165                1170

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1175                1180                1185

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1190                1195                1200

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1205                1210                1215

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1220                1225                1230

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1235                1240                1245

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1250                1255                1260

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1265                1270                1275

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1280                1285                1290

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1295                1300                1305

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1310                1315                1320

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Gly Ala  Pro Gly Ser
    1325                1330                1335
```

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
1340 1345 1350

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
1355 1360 1365

Ser Glu Thr Pro Ala Ser Ser Glu Trp Leu Gln Val Asp Phe Gln
1370 1375 1380

Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser
1385 1390 1395

Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser
1400 1405 1410

Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val
1415 1420 1425

Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn
1430 1435 1440

Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro
1445 1450 1455

Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly
1460 1465 1470

Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu
1475 1480 1485

Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala
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Thr Gly His His His His His His
1505 1510

<210> 676
<211> 4536
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

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ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

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atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

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gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

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aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

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<210> 677
<211> 1511
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 677

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

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His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480
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Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750
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Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020
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Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
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Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1040                1045                1050

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1055                1060                1065

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1070                1075                1080

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1085                1090                1095

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1100                1105                1110

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1115                1120                1125

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1130                1135                1140

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1145                1150                1155

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1160                1165                1170

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1175                1180                1185

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1190                1195                1200

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1205                1210                1215

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1220                1225                1230

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1235                1240                1245

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1250                1255                1260

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1265                1270                1275
```

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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1280                 1285                 1290

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1295                 1300                 1305

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1310                 1315                 1320

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1325                 1330                 1335

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1340                 1345                 1350

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1355                 1360                 1365

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1370                 1375                 1380

Gly Ala  Pro Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1385                 1390                 1395

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    1400                 1405                 1410

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Ala Ser Ser  Gly Lys Val
    1415                 1420                 1425

Lys Val  Phe Gln Gly Asn Gln  Asp Ser Phe Thr Pro  Val Val Asn
    1430                 1435                 1440

Ser Leu  Asp Pro Pro Leu Leu  Thr Arg Tyr Leu Arg  Ile His Pro
    1445                 1450                 1455

Gln Ser  Trp Val His Gln Ile  Ala Leu Arg Met Glu  Val Leu Gly
    1460                 1465                 1470

Cys Glu  Ala Gln Asp Leu Tyr  Gly Ala Gly Ser Pro  Gly Ala Glu
    1475                 1480                 1485

Thr Ala  Glu Gln Lys Leu Ile  Ser Glu Glu Asp Leu  Ser Pro Ala
    1490                 1495                 1500

Thr Gly  His His His His His  His
    1505                 1510
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<210> 678
<211> 4536
<212> DNA

<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

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tgggcttatt tctctgatgt tgacctggaa aaagatgtgc actcaggcct gattggaccc 2880

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ctgttggcac caatgattat tcacggcatc aagacccagg gtgcccgtca gaagttctcc 3600

agcctctaca tctctcagtt tatcatcatg tatagtcttg atgggaagaa gtggcagact 3660

tatcgaggaa attccactgg aaccttaatg gtcttctttg gcaatgtgga ttcatctggg 3720

ataaaacaca atatttttaa ccctccaatt attgctcgat acatccgttt gcacccaact 3780

cattatagca ttcgcagcac tcttcgcatg gagttgatgg gctgtgattt aaatagttgc 3840

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tactttacca atatgtttgc cacctggtct ccttcaaaag ctcgacttca cctccaaggg 3960

aggagtaatg cctggagacc tcaggtgaat aatccaaaag agtggctgca agtggacttc 4020

cagaagacaa tgaaagtcac aggagtaact actcagggag taaaatctct gcttaccagc 4080

atgtatgtga aggagttcct catctccagc agtcaagatg gccatcagtg gactctcttt 4140

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agcgaaagcg ctacgcctga gagtggccct ggctctgagc cagccacctc cggctctgaa 4260

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<210> 679
<211> 1762
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 679

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

```
Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415
```

```
Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685
```

```
Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960
```

```
Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1025                1030                1035

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1040                1045                1050

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1055                1060                1065

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1070                1075                1080

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1085                1090                1095

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1100                1105                1110

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1115                1120                1125

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1130                1135                1140

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1145                1150                1155

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1160                1165                1170

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1175                1180                1185

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1190                1195                1200

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1205                1210                1215
```

```
Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1220                 1225                 1230

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1235                 1240                 1245

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1250                 1255                 1260

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1265                 1270                 1275

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1280                 1285                 1290

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1295                 1300                 1305

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1310                 1315                 1320

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1325                 1330                 1335

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1340                 1345                 1350

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1355                 1360                 1365

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1370                 1375                 1380

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1385                 1390                 1395

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1400                 1405                 1410

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1415                 1420                 1425

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1430                 1435                 1440

Gly Ala  Glu Thr Ala Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1445                 1450                 1455

Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly
    1460                 1465                 1470
```

```
Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro
    1475                 1480                 1485

Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr
    1490                 1495                 1500

Pro Glu  Ser Gly Pro Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    1505                 1510                 1515

Ala Pro  Gly Ser Pro Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly
    1520                 1525                 1530

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro
    1535                 1540                 1545

Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr
    1550                 1555                 1560

Pro Glu  Ser Gly Pro Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr
    1565                 1570                 1575

Glu Glu  Gly Ser Pro Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly
    1580                 1585                 1590

Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Glu
    1595                 1600                 1605

Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr Ser Glu  Ser Ala Thr
    1610                 1615                 1620

Pro Glu  Ser Gly Pro Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1625                 1630                 1635

Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly
    1640                 1645                 1650

Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr Pro Gly  Ser Pro Ala
    1655                 1660                 1665

Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr Ser Thr  Glu Pro Ser
    1670                 1675                 1680

Glu Gly  Ser Ala Pro Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    1685                 1690                 1695

Ala Pro  Gly Ser Glu Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly
    1700                 1705                 1710

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Thr
    1715                 1720                 1725
```

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Glu Thr Ala Glu Gln
1730 1735 1740

Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala Thr Gly His His
1745 1750 1755

His His His His
1760

<210> 680
<211> 5289
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

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gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

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<210> 681

<211> 1762
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 681

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240
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```
Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510
```

```
Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780
```

```
Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1025                1030                1035

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1040                1045                1050
```

```
Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1055                1060                1065

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1070                1075                1080

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1085                1090                1095

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1100                1105                1110

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1115                1120                1125

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1130                1135                1140

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1145                1150                1155

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1160                1165                1170

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1175                1180                1185

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1190                1195                1200

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1205                1210                1215

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1220                1225                1230

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1235                1240                1245

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1250                1255                1260

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1265                1270                1275

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1280                1285                1290

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1295                1300                1305
```

```
Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1310                1315                1320

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1325                1330                1335

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1340                1345                1350

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1355                1360                1365

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1370                1375                1380

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1385                1390                1395

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1400                1405                1410

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1415                1420                1425

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1430                1435                1440

Gly Ala  Glu Thr Ala Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
    1445                1450                1455

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1460                1465                1470

Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser Ser Pro  Gly Ser Ser
    1475                1480                1485

Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Thr Pro  Gly Ser Gly
    1490                1495                1500

Thr Ala  Ser Ser Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    1505                1510                1515

Gly Ser  Pro Gly Thr Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro
    1520                1525                1530

Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser
    1535                1540                1545

Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala
    1550                1555                1560
```

Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
1565 1570 1575

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
1580 1585 1590

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser
1595 1600 1605

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala
1610 1615 1620

Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
1625 1630 1635

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
1640 1645 1650

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
1655 1660 1665

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala
1670 1675 1680

Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
1685 1690 1695

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
1700 1705 1710

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser
1715 1720 1725

Thr Pro Ser Gly Ala Thr Gly Ser Gly Ala Glu Thr Ala Glu Gln
1730 1735 1740

Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala Thr Gly His His
1745 1750 1755

His His His His
1760

<210> 682
<211> 5289
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 682
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ctcggtgagc tgcctgtgga cgcaagattt cctcctagag tgccaaaatc ttttccattc 120

aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

agagatctag cttcaggact cattggccct ctcctcatct gctacaaaga atctgtagat 1680

caaagaggaa accagataat gtcagacaag aggaatgtca tcctgttttc tgtatttgat 1740

gagaaccgaa gctggtacct cacagagaat atacaacgct ttctccccaa tccagctgga 1800

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gtttttgata gtttgcagtt gtcagtttgt ttgcatgagg tggcatactg gtacattcta 1920

agcattggag cacagactga cttcctttct gtcttcttct ctggatatac cttcaaacac 1980

aaaatggtct atgaagacac actcacccta ttcccattct caggagaaac tgtcttcatg 2040

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ggcatgaccg ccttactgaa ggtttctagt tgtgacaaga acactggtga ttattacgag 2160

gacagttatg aagatatttc agcatacttg ctgagtaaaa acaatgccat tgaaccaaga 2220

agcttctctc aaaacccacc agtcttgaaa cgccatcaac gggaaataac tcgtactact 2280

cttcagtcag atcaagagga aatcgattat gatgatacca tatcagttga aatgaagaag 2340

gaagattttg acatttatga tgaggatgaa aatcagagcc cccgcagctt tcaaaagaaa 2400

acacgacact attttattgc tgcagtggag aggctctggg attatgggat gagtagctcc 2460

ccacatgttc taagaaacag ggctcagagt ggcagtgtcc ctcagttcaa gaaagttgtt 2520

ttccaggaat ttactgatgg ctcctttact cagcccttat accgtggaga actaaatgaa 2580

catttgggac tcctggggcc atatataaga gcagaagttg aagataatat catggtaact 2640

ttcagaaatc aggcctctcg tccctattcc ttctattcta gccttatttc ttatgaggaa 2700

gatcagaggc aaggagcaga acctagaaaa aactttgtca agcctaatga aaccaaaact 2760

tacttttgga aagtgcaaca tcatatggca cccactaaag atgagtttga ctgcaaagcc 2820

tgggcttatt tctctgatgt tgacctggaa aaagatgtgc actcaggcct gattggaccc 2880

cttctggtct gccacactaa cacactgaac cctgctcatg ggagacaagt gacagtacag 2940

gaatttgctc tgtttttcac catctttgat gagaccaaaa gctggtactt cactgaaaat 3000

atggaaagaa actgcagggc tccctgcaat atccagatgg aagatcccac ttttaaagag 3060

aattatcgct tccatgcaat caatggctac ataatggata cactacctgg cttagtaatg 3120

gctcaggatc aaaggattcg atggtatctg ctcagcatgg gcagcaatga aaacatccat 3180

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ataaaacaca atatttttaa ccctccaatt attgctcgat acatccgttt gcacccaact 3780

cattatagca ttcgcagcac tcttcgcatg gagttgatgg gctgtgattt aaatagttgc 3840

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tactttacca atatgtttgc cacctggtct ccttcaaaag ctcgacttca cctccaaggg 3960

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cagaagacaa tgaaagtcac aggagtaact actcagggag taaaatctct gcttaccagc 4080

atgtatgtga aggagttcct catctccagc agtcaagatg gccatcagtg gactctcttt 4140

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gcgtcgtcga gcccagggtc gtccacccca agcggggcta ctgggtcccc gggtagctcg 4860

ccgagcgcat caacgggaac cggaccgggc agctcccccct cagcgtcgac agggacaggc 4920

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gcgtcaagct cgcccggaag ctccacgcct tcgggggcaa ccgggtcagg ggccgaaacg 5220

gccgaacaaa aactcatctc agaagaggat ctgtcacctg caaccggtca tcatcaccat 5280

caccattga 5289

<210> 683
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 683
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

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Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335
```

```
Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
                405                 410                 415

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            420                 425                 430

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
        435                 440                 445

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    450                 455                 460

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
465                 470                 475                 480

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
                485                 490                 495

Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            500                 505                 510

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
        515                 520                 525

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
    530                 535                 540

Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605
```

```
Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880
```

```
Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140
```

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Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395
```

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1400 1405 1410

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1415 1420 1425

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1430 1435 1440

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1445 1450 1455

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1460 1465 1470

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1475 1480 1485

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1490 1495 1500

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1505 1510 1515

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1520 1525 1530

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1580 1585 1590

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1595 1600 1605

Ser Pro Ala Thr Gly His His His His His His
1610 1615

<210> 684
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 684
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ctcggtgagc tgcctgtgga cgcaagattt cctcctagag tgccaaaatc ttttccattc 120

aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg gcgcgccaac cagtacggag ccgtccgagg ggagcgcacc aggaagcccg 1260

gctgggagcc cgacttctac cgaagagggt acatctaccg aaccaagtga aggttcagca 1320

ccaggcacct caacagaacc ctctgagggc tcggcgcctg gtacaagtga gtccgccacc 1380

ccagaatccg ggcctgggac aagcacagaa ccttcggaag ggagtgcccc tggaacatcc 1440

gaatcggcaa ccccagaatc agggccagga tctgagcccg cgacttcggg ctccgagacg 1500

cctgggacat ccaccgagcc ctccgaagga tcagccccag gcaccagcac ggagccctct 1560

gagggaagcg cacctggtac cagcgaaagc gcaactcccg aatcaggtcc cggtacgagc 1620

gagtcggcga ccccggagag cgggccaggt gcctcgagcg acagaagtta taaaagtcaa 1680

tatttgaaca atggccctca gcggattggt aggaagtaca aaaaagtccg atttatggca 1740

tacacagatg aaacctttaa gactcgtgaa gctattcagc atgaatcagg aatcttggga 1800

cctttacttt atggggaagt tggagacaca ctgttgatta tatttaagaa tcaagcaagc 1860

agaccatata acatctaccc tcacggaatc actgatgtcc gtcctttgta ttcaaggaga 1920

ttaccaaaag gtgtaaaaca tttgaaggat tttccaattc tgccaggaga aatattcaaa 1980

tataaatgga cagtgactgt agaagatggg ccaactaaat cagatcctcg gtgcctgacc 2040

cgctattact ctagtttcgt taatatggag agagatctag cttcaggact cattggccct 2100

ctcctcatct gctacaaaga atctgtagat caaagaggaa accagataat gtcagacaag 2160

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tccaacatca tgcacagcat caatggctat gtttttgata gtttgcagtt gtcagtttgt 2340

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ttcccattct caggagaaac tgtcttcatg tcgatggaaa acccaggtct atggattctg 2520

gggtgccaca actcagactt tcggaacaga ggcatgaccg ccttactgaa ggtttctagt 2580

tgtgacaaga acactggtga ttattacgag gacagttatg aagatatttc agcatacttg 2640

ctgagtaaaa acaatgccat tgaaccaaga agcttctctc aaaacccacc agtcttgaaa 2700

cgccatcaac gggaaataac tcgtactact cttcagtcag atcaagagga aatcgattat 2760

gatgatacca tatcagttga aatgaagaag gaagattttg acatttatga tgaggatgaa 2820

aatcagagcc cccgcagctt tcaaaagaaa acacgacact attttattgc tgcagtggag 2880

aggctctggg attatgggat gagtagctcc ccacatgttc taagaaacag ggctcagagt 2940

ggcagtgtcc ctcagttcaa gaaagttgtt ttccaggaat ttactgatgg ctcctttact 3000

cagcccttat accgtggaga actaaatgaa catttgggac tcctggggcc atatataaga 3060

gcagaagttg aagataatat catggtaact ttcagaaatc aggcctctcg tccctattcc 3120

ttctattcta gccttatttc ttatgaggaa gatcagaggc aaggagcaga acctagaaaa 3180

aactttgtca agcctaatga aaccaaaact tacttttgga aagtgcaaca tcatatggca 3240

cccactaaag atgagtttga ctgcaaagcc tgggcttatt tctctgatgt tgacctggaa 3300

aaagatgtgc actcaggcct gattggaccc cttctggtct gccacactaa cacactgaac 3360

cctgctcatg ggagacaagt gacagtacag gaatttgctc tgtttttcac catctttgat 3420

gagaccaaaa gctggtactt cactgaaaat atggaaagaa actgcagggc tccctgcaat 3480

atccagatgg aagatcccac ttttaaagag aattatcgct tccatgcaat caatggctac 3540

ataatggata cactacctgg cttagtaatg gctcaggatc aaaggattcg atggtatctg 3600

ctcagcatgg gcagcaatga aaacatccat tctattcatt tcagtggaca tgtgttcact 3660

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ggaaatcaag actccttcac acctgtggtg aactctctag acccaccgtt actgactcgc 4680

taccttcgaa ttcaccccca gagttgggtg caccagattg ccctgaggat ggaggttctg 4740

ggctgcgagg cacaggacct ctacggcgcc ggatcacctg gggccgaaac ggccgaacaa 4800

aaactcatct cagaagagga tctgtcacct gcaaccggtc atcatcacca tcaccattga 4860

<210> 685
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 685
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val

115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu

385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Ala Pro Thr Ser Thr Glu Pro Ser
595 600 605

Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
610 615 620

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
625 630 635 640

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
645 650 655

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala

660 665 670

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
675 680 685

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser
690 695 700

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
705 710 715 720

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
725 730 735

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
835 840 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
885 890 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
900 905 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
915 920 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro

930 935 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945 950 955 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
965 970 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
980 985 990

Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
995 1000 1005

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1010 1015 1020

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1025 1030 1035

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1040 1045 1050

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1055 1060 1065

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1070 1075 1080

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1085 1090 1095

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1100 1105 1110

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1115 1120 1125

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1130 1135 1140

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1145 1150 1155

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1160 1165 1170

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1175 1180 1185

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met

1190 1195 1200

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1205 1210 1215

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1220 1225 1230

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1235 1240 1245

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1250 1255 1260

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1265 1270 1275

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1280 1285 1290

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1295 1300 1305

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1310 1315 1320

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1325 1330 1335

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1340 1345 1350

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1355 1360 1365

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1370 1375 1380

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1385 1390 1395

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1400 1405 1410

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1415 1420 1425

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1430 1435 1440

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr

1445 1450 1455

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1460 1465 1470

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1475 1480 1485

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1490 1495 1500

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1505 1510 1515

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1520 1525 1530

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1580 1585 1590

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1595 1600 1605

Ser Pro Ala Thr Gly His His His His His His
1610 1615

<210> 686
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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ctcggtgagc tgcctgtgga cgcaagattt cctcctagag tgccaaaatc ttttccattc 120

aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420
ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480
catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540
gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600
gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660
gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720
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atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960
gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020
gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

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Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
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His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
                405                 410                 415

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser
            420                 425                 430

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly
        435                 440                 445
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Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
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Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala
465                 470                 475                 480

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly
                485                 490                 495

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
            500                 505                 510

Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala
        515                 520                 525

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
    530                 535                 540

Ser Ser Thr Gly Ser Pro Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720
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Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
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Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990
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Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
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Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
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Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
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Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
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Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
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Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
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Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
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Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
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Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
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Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
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Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
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Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
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Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
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Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
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Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
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Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
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Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
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Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
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Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
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Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
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Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
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Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1490                1495                1500
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Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
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Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1520 1525 1530

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
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gagttgatgg gctgtgattt aaatagttgc agcatgccat tgggaatgga gagtaaagca 4320

atatcagatg cacagattac tgcttcatcc tactttacca atatgtttgc cacctggtct 4380

ccttcaaaag ctcgacttca cctccaaggg aggagtaatg cctggagacc tcaggtgaat 4440

aatccaaaag agtggctgca agtggacttc cagaagacaa tgaaagtcac aggagtaact 4500

actcagggag taaaatctct gcttaccagc atgtatgtga aggagttcct catctccagc 4560

agtcaagatg gccatcagtg gactctcttt tttcagaatg gcaaagtaaa ggtttttcag 4620

ggaaatcaag actccttcac acctgtggtg aactctctag acccaccgtt actgactcgc 4680

taccttcgaa ttcaccccca gagttgggtg caccagattg ccctgaggat ggaggttctg 4740

ggctgcgagg cacaggacct ctacggcgcc ggatcacctg gggccgaaac ggccgaacaa 4800

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<210> 689
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 689
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

```
Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495
```

```
His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Ala Pro Gly Ala Ser Pro Gly Thr
        595                 600                 605

Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr
    610                 615                 620

Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr
625                 630                 635                 640

Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser
                645                 650                 655

Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr
            660                 665                 670

Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr
        675                 680                 685

Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser
    690                 695                 700

Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
705                 710                 715                 720

Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
                725                 730                 735

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765
```

```
Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035
```

```
Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290
```

```
Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                1435                1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                1450                1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                1465                1470

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                1480                1485

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1490                1495                1500

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                1510                1515

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1520                1525                1530

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1535                1540                1545
```

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1580 1585 1590

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1595 1600 1605

Ser Pro Ala Thr Gly His His His His His His
1610 1615

<210> 690
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

agagatctag cttcaggact cattggccct ctcctcatct gctacaaaga atctgtagat 1680

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gagaaccgaa gctggtacct cacagagaat atacaacgct ttctccccaa tccagctggc 1800

gcgccaggtg cctcgccggg aacatcatca actggttcac ccgggtcatc cccctcggcc 1860

tcaaccggga cgggtcccgg ctcatccccc agcgccagca ctggaacagg tcctggcact 1920

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tcacctggct cgtcaccttc ggcgtcaact ggaacggggc caggggcctc acccggaacg 2040

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gggtgccaca actcagactt tcggaacaga ggcatgaccg ccttactgaa ggtttctagt 2580

tgtgacaaga acactggtga ttattacgag gacagttatg aagatatttc agcatacttg 2640

ctgagtaaaa acaatgccat tgaaccaaga agcttctctc aaaacccacc agtcttgaaa 2700

cgccatcaac gggaaataac tcgtactact cttcagtcag atcaagagga aatcgattat 2760

gatgatacca tatcagttga aatgaagaag gaagattttg acatttatga tgaggatgaa 2820

aatcagagcc cccgcagctt tcaaaagaaa acacgacact attttattgc tgcagtggag 2880

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cagcccttat accgtggaga actaaatgaa catttgggac tcctggggcc atatataaga 3060

gcagaagttg aagataatat catggtaact ttcagaaatc aggcctctcg tccctattcc 3120

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aactttgtca agcctaatga aaccaaaact tacttttgga aagtgcaaca tcatatggca 3240

cccactaaag atgagtttga ctgcaaagcc tgggcttatt tctctgatgt tgacctggaa 3300

aaagatgtgc actcaggcct gattggaccc cttctggtct gccacactaa cacactgaac 3360

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gagaccaaaa gctggtactt cactgaaaat atggaaagaa actgcagggc tccctgcaat 3480

atccagatgg aagatcccac ttttaaagag aattatcgct tccatgcaat caatggctac 3540

ataatggata cactacctgg cttagtaatg gctcaggatc aaaggattcg atggtatctg 3600

ctcagcatgg gcagcaatga aaacatccat tctattcatt tcagtggaca tgtgttcact 3660

gtacgaaaaa aagaggagta taaaatggca ctgtacaatc tctatccagg tgtttttgag 3720

acagtggaaa tgttaccatc caaagctgga atttggcggg tggaatgcct tattggcgag 3780

catctacatg ctgggatgag cacacttttt ctggtgtaca gcaataagtg tcagactccc 3840

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gtcttctttg gcaatgtgga ttcatctggg ataaaacaca atatttttaa ccctccaatt 4200

attgctcgat acatccgttt gcacccaact cattatagca ttcgcagcac tcttcgcatg 4260

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agtcaagatg gccatcagtg gactctcttt tttcagaatg gcaaagtaaa ggtttttcag 4620

ggaaatcaag actccttcac acctgtggtg aactctctag acccaccgtt actgactcgc 4680

taccttcgaa ttcaccccca gagttgggtg caccagattg ccctgaggat ggaggttctg 4740

ggctgcgagg cacaggacct ctacggcgcc ggatcacctg gggccgaaac ggccgaacaa 4800

aaactcatct cagaagagga tctgtcacct gcaaccggtc atcatcacca tcaccattga 4860

<210> 691
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 691

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Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1               5                  10                  15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270
```

```
Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540
```

```
Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
            740                 745                 750

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
        755                 760                 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
    770                 775                 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785                 790                 795                 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                805                 810                 815
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Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            820                 825                 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
        835                 840                 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    850                 855                 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865                 870                 875                 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080
```

```
Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335
```

```
Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                1435                1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                1450                1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                1465                1470

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                1480                1485

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1490                1495                1500

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                1510                1515

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1520                1525                1530

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1535                1540                1545

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1550                1555                1560

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1565                1570                1575

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1580                1585                1590
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Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1595 1600 1605

Ser Pro Ala Thr Gly His His His His His His
1610 1615

<210> 692
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 692
gccaccagaa gatactacct gggtgcagtg gaactgtcat gggactatat gcaaagtgat 60

ctcggtgagc tgcctgtgga cgcaagattt cctcctagag tgccaaaatc ttttccattc 120

aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

agagatctag cttcaggact cattggccct ctcctcatct gctacaaaga atctgtagat 1680

caaagaggaa accagataat gtcagacaag aggaatgtca tcctgttttc tgtatttgat 1740

gagaaccgaa gctggtacct cacagagaat atacaacgct ttctccccaa tccagctgga 1800

gtgcagcttg aggatccaga gttccaagcc tccaacatca tgcacagcat caatggctat 1860

gtttttgata gtttgcagtt gtcagtttgt ttgcatgagg tggcatactg gtacattcta 1920

agcattggag cacagactga cttccttttct gtcttcttct ctggatatac cttcaaacac 1980

aaaatggtct atgaagacac actcacccta ttcccattct caggagaaac tgtcttcatg 2040

tcgatggaaa acccaggtct atggattctg gggtgccaca actcagactt tcggaacaga 2100

ggcatgaccg ccttactgaa ggtttctagt tgtgacaaga acactggtga ttattacgag 2160

gacagttatg aagatatttc agcatacttg ctgagtaaaa acaatgccat tgaaccaaga 2220

agcttctctc aaaacggcgc gccaagtccc gctggaagcc caactagcac cgaagagggg 2280

acctcagagt ccgccacccc cgagtccggc cctggctctg agcctgccac tagcggctcc 2340

gagactcctg gcacatccga aagcgctaca cccgagagtg gacccggcac ctctaccgag 2400

cccagtgagg gctccgcccc tggaacaagc accgagccca gcgaaggcag cgccccaggg 2460

acctccacag agcccagtga aggcagtgct cctggcacca gcaccgaacc aagcgagggc 2520

tctgcacccg ggacctccac cgagccaagc gaaggctctg cccctggcac ttccaccgag 2580

cccagcgaag gcagcgcccc tgggagcccc gctggctctc ccaccagcac tgaggagggc 2640

acatctaccg aaccaagtga aggctctgca ccaggtgcct cgagcccacc agtcttgaaa 2700

cgccatcaac gggaaataac tcgtactact cttcagtcag atcaagagga aatcgattat 2760

gatgatacca tatcagttga aatgaagaag gaagattttg acatttatga tgaggatgaa 2820

aatcagagcc cccgcagctt tcaaaagaaa acacgacact attttattgc tgcagtggag 2880

aggctctggg attatgggat gagtagctcc ccacatgttc taagaaacag ggctcagagt 2940

ggcagtgtcc ctcagttcaa gaaagttgtt ttccaggaat ttactgatgg ctcctttact 3000

cagcccttat accgtggaga actaaatgaa catttgggac tcctggggcc atatataaga 3060

gcagaagttg aagataatat catggtaact ttcagaaatc aggcctctcg tccctattcc 3120

ttctattcta gccttatttc ttatgaggaa gatcagaggc aaggagcaga acctagaaaa 3180

aactttgtca agcctaatga aaccaaaact tacttttgga aagtgcaaca tcatatggca 3240

cccactaaag atgagtttga ctgcaaagcc tgggcttatt tctctgatgt tgacctggaa 3300

aaagatgtgc actcaggcct gattggaccc cttctggtct gccacactaa cacactgaac 3360

cctgctcatg ggagacaagt gacagtacag gaatttgctc tgtttttcac catctttgat 3420

gagaccaaaa gctggtactt cactgaaaat atggaaagaa actgcagggc tccctgcaat 3480

atccagatgg aagatcccac ttttaaagag aattatcgct tccatgcaat caatggctac 3540

ataatggata cactacctgg cttagtaatg gctcaggatc aaaggattcg atggtatctg 3600

ctcagcatgg gcagcaatga aaacatccat tctattcatt tcagtggaca tgtgttcact 3660

gtacgaaaaa aagaggagta taaaatggca ctgtacaatc tctatccagg tgtttttgag 3720

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aagacccagg gtgcccgtca gaagttctcc agcctctaca tctctcagtt tatcatcatg 4080

tatagtcttg atgggaagaa gtggcagact tatcgaggaa attccactgg aaccttaatg 4140

gtcttctttg gcaatgtgga ttcatctggg ataaaacaca atatttttaa ccctccaatt 4200

attgctcgat acatccgttt gcacccaact cattatagca ttcgcagcac tcttcgcatg 4260

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ggctgcgagg cacaggacct ctacggcgcc ggatcacctg gggccgaaac ggccgaacaa 4800

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<210> 693
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 693
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro

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    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
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325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe

595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
740 745 750

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
755 760 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
770 775 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785 790 795 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
805 810 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
820 825 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
835 840 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
850 855 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly

865 870 875 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
885 890 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
900 905 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
915 920 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
930 935 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945 950 955 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
965 970 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
980 985 990

Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
995 1000 1005

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1010 1015 1020

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1025 1030 1035

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1040 1045 1050

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1055 1060 1065

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1070 1075 1080

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1085 1090 1095

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1100 1105 1110

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1115 1120 1125

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys

1130 1135 1140

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1145 1150 1155

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1160 1165 1170

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1175 1180 1185

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1190 1195 1200

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1205 1210 1215

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1220 1225 1230

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1235 1240 1245

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1250 1255 1260

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1265 1270 1275

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1280 1285 1290

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1295 1300 1305

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1310 1315 1320

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1325 1330 1335

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1340 1345 1350

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1355 1360 1365

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1370 1375 1380

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro

1385 1390 1395

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1400 1405 1410

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1415 1420 1425

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1430 1435 1440

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1445 1450 1455

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1460 1465 1470

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1475 1480 1485

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1490 1495 1500

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1505 1510 1515

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1520 1525 1530

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1580 1585 1590

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1595 1600 1605

Ser Pro Ala Thr Gly His His His His His His
1610 1615

<210> 694
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 694
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

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Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380
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Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
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Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655
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Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Thr Pro Gly
            740                 745                 750

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
        755                 760                 765

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    770                 775                 780

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
785                 790                 795                 800

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
                805                 810                 815

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            820                 825                 830

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
        835                 840                 845

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
    850                 855                 860

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
865                 870                 875                 880

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
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Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
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Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
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Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
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Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
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Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
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Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
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Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
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Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
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Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
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Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
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Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
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Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
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Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
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Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
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Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
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Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
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Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
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Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
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Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
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Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
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Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                1435                1440
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Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
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Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
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Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
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Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
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Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
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Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
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Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
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Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
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cccactaaag atgagtttga ctgcaaagcc tgggcttatt tctctgatgt tgacctggaa 3300

aaagatgtgc actcaggcct gattggaccc cttctggtct gccacactaa cacactgaac 3360

cctgctcatg ggagacaagt gacagtacag gaatttgctc tgtttttcac catctttgat 3420

gagaccaaaa gctggtactt cactgaaaat atggaaagaa actgcagggc tccctgcaat 3480

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ataatggata cactacctgg cttagtaatg gctcaggatc aaaggattcg atggtatctg 3600

ctcagcatgg gcagcaatga aaacatccat tctattcatt tcagtggaca tgtgttcact 3660

gtacgaaaaa aagaggagta taaaatggca ctgtacaatc tctatccagg tgtttttgag 3720

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catctacatg ctgggatgag cacacttttt ctggtgtaca gcaataagtg tcagactccc 3840

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ggctgcgagg cacaggacct ctacggcgcc ggatcacctg gggccgaaac ggccgaacaa 4800

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<210> 697
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 697

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

```
His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430
```

```
Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700
```

```
Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Thr Pro Gly
            740                 745                 750

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
        755                 760                 765

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    770                 775                 780

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
785                 790                 795                 800

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
                805                 810                 815

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            820                 825                 830

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
        835                 840                 845

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
    850                 855                 860

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
865                 870                 875                 880

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975
```

```
Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230
```

```
Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                1435                1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                1450                1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                1465                1470

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                1480                1485
```

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1490 1495 1500

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1505 1510 1515

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1520 1525 1530

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1580 1585 1590

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1595 1600 1605

Ser Pro Ala Thr Gly His His His His His His
1610 1615

<210> 698
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

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atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

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gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

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<210> 699
<211> 1622
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 699

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Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205
```

```
His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480
```

```
Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750
```

```
Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Gly Ala Pro
            820                 825                 830

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
        835                 840                 845

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
    850                 855                 860

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
865                 870                 875                 880

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
                885                 890                 895

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
            900                 905                 910

Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
        915                 920                 925

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
    930                 935                 940

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
945                 950                 955                 960

Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
                965                 970                 975

Ala Ser Ser Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val
            980                 985                 990

Val Phe Gln Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
        995                 1000                1005

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1010                1015                1020
```

```
Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1025                1030                1035

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1040                1045                1050

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1055                1060                1065

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1070                1075                1080

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1085                1090                1095

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1100                1105                1110

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1115                1120                1125

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1130                1135                1140

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1145                1150                1155

Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1160                1165                1170

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1175                1180                1185

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1190                1195                1200

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1205                1210                1215

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1220                1225                1230

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1235                1240                1245

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1250                1255                1260

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1265                1270                1275
```

```
Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1280                1285                1290

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1295                1300                1305

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1310                1315                1320

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1325                1330                1335

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1340                1345                1350

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1355                1360                1365

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1370                1375                1380

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1385                1390                1395

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1400                1405                1410

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1415                1420                1425

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1430                1435                1440

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1445                1450                1455

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1460                1465                1470

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1475                1480                1485

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1490                1495                1500

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1505                1510                1515

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1520                1525                1530
```

Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser
1535 1540 1545

Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg
1550 1555 1560

Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu
1565 1570 1575

Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala
1580 1585 1590

Gly Ser Pro Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu
1595 1600 1605

Glu Asp Leu Ser Pro Ala Thr Gly His His His His His His
1610 1615 1620

<210> 700
<211> 4869
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 700
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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

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aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

agagatctag cttcaggact cattggccct ctcctcatct gctacaaaga atctgtagat 1680

caaagaggaa accagataat gtcagacaag aggaatgtca tcctgttttc tgtatttgat 1740

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ggcatgaccg ccttactgaa ggtttctagt tgtgacaaga acactggtga ttattacgag 2160

gacagttatg aagatatttc agcatacttg ctgagtaaaa acaatgccat tgaaccaaga 2220

agcttctctc aaaacccacc agtcttgaaa cgccatcaac gggaaataac tcgtactact 2280

cttcagtcag atcaagagga aatcgattat gatgatacca tatcagttga aatgaagaag 2340

gaagattttg acatttatga tgaggatgaa aatcagagcc cccgcagctt tcaaaagaaa 2400

acacgacact attttattgc tgcagtggag aggctctggg attatgggat gagtagctcc 2460

ccacatgttc taagaaacag ggctcagggc gcgccaacgt ccgaaagtgc tacccctgag 2520

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catatggcac ccactaaaga tgagtttgac tgcaaagcct gggcttattt ctctgatgtt 3300

gacctggaaa aagatgtgca ctcaggcctg attggacccc ttctggtctg ccacactaac 3360

acactgaacc ctgctcatgg gagacaagtg acagtacagg aatttgctct gtttttcacc 3420

atctttgatg agaccaaaag ctggtacttc actgaaaata tggaaagaaa ctgcagggct 3480

ccctgcaata tccagatgga agatcccact tttaaagaga attatcgctt ccatgcaatc 3540

aatggctaca taatggatac actacctggc ttagtaatgg ctcaggatca aaggattcga 3600

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gtttttgaga cagtggaaat gttaccatcc aaagctggaa tttggcgggt ggaatgcctt 3780

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cagactcccc tgggaatggc ttctggacac attagagatt ttcagattac agcttcagga 3900

caatatggac agtgggcccc aaagctggcc agacttcatt attccggatc aatcaatgcc 3960

tggagcacca aggagccctt ttcttggatc aaggtggatc tgttggcacc aatgattatt 4020

cacggcatca agacccaggg tgcccgtcag aagttctcca gcctctacat ctctcagttt 4080

atcatcatgt atagtcttga tgggaagaag tggcagactt atcgaggaaa ttccactgga 4140

accttaatgg tcttctttgg caatgtggat tcatctggga taaaacacaa tatttttaac 4200

cctccaatta ttgctcgata catccgtttg cacccaactc attatagcat tcgcagcact 4260

cttcgcatgg agttgatggg ctgtgattta aatagttgca gcatgccatt gggaatggag 4320

agtaaagcaa tatcagatgc acagattact gcttcatcct actttaccaa tatgtttgcc 4380

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atctccagca gtcaagatgg ccatcagtgg actctctttt ttcagaatgg caaagtaaag 4620

gtttttcagg gaaatcaaga ctccttcaca cctgtggtga actctctaga cccaccgtta 4680

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gaggttctgg gctgcgaggc acaggacctc tacggcgccg gatcacctgg ggccgaaacg 4800

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caccattga 4869

<210> 701
<211> 1622
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 701

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Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255
```

```
Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525
```

```
Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800
```

```
Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Gly Ala Pro
            820                 825                 830

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
        835                 840                 845

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
    850                 855                 860

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
865                 870                 875                 880

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
                885                 890                 895

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
            900                 905                 910

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
        915                 920                 925

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
    930                 935                 940

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
945                 950                 955                 960

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                965                 970                 975

Ala Ser Ser Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val
            980                 985                 990

Val Phe Gln Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
        995                 1000                1005

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1010                1015                1020

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1025                1030                1035

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1040                1045                1050

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1055                1060                1065
```

```
Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1070                1075                1080

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1085                1090                1095

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1100                1105                1110

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1115                1120                1125

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1130                1135                1140

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1145                1150                1155

Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1160                1165                1170

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1175                1180                1185

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1190                1195                1200

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1205                1210                1215

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1220                1225                1230

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1235                1240                1245

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1250                1255                1260

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1265                1270                1275

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1280                1285                1290

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1295                1300                1305

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1310                1315                1320
```

```
Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1325                1330                1335

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1340                1345                1350

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1355                1360                1365

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1370                1375                1380

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1385                1390                1395

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1400                1405                1410

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1415                1420                1425

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1430                1435                1440

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1445                1450                1455

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1460                1465                1470

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1475                1480                1485

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1490                1495                1500

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1505                1510                1515

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1520                1525                1530

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1535                1540                1545

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    1550                1555                1560

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    1565                1570                1575
```

Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala
1580 1585 1590

Gly Ser Pro Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu
1595 1600 1605

Glu Asp Leu Ser Pro Ala Thr Gly His His His His His His
1610 1615 1620

<210> 702
<211> 4869
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

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aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

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caaagaggaa accagataat gtcagacaag aggaatgtca tcctgttttc tgtatttgat 1740

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agcttctctc aaaacccacc agtcttgaaa cgccatcaac gggaaataac tcgtactact 2280

cttcagtcag atcaagagga aatcgattat gatgatacca tatcagttga aatgaagaag 2340

gaagattttg acatttatga tgaggatgaa aatcagagcc cccgcagctt tcaaaagaaa 2400

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caatatggac agtgggcccc aaagctggcc agacttcatt attccggatc aatcaatgcc 3960

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caccattga 4869

<210> 703
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300
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Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575
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Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845
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Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1025                1030                1035

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1040                1045                1050

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1055                1060                1065

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1070                1075                1080

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1085                1090                1095

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1100                1105                1110
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```
Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1115                1120                1125

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1130                1135                1140

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1145                1150                1155

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1160                1165                1170

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1175                1180                1185

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1190                1195                1200

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1205                1210                1215

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1220                1225                1230

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1235                1240                1245

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1250                1255                1260

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Gly
    1265                1270                1275

Ala Pro  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1280                1285                1290

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    1295                1300                1305

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1310                1315                1320

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1325                1330                1335

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    1340                1345                1350

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    1355                1360                1365
```

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Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1370                1375                1380

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1385                1390                1395

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser
    1400                1405                1410

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ala  Ser Ser Asn
    1415                1420                1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                1435                1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                1450                1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                1465                1470

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                1480                1485

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1490                1495                1500

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                1510                1515

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1520                1525                1530

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1535                1540                1545

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1550                1555                1560

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1565                1570                1575

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1580                1585                1590

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
    1595                1600                1605

Ser Pro  Ala Thr Gly His His  His His His His His
    1610                1615
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<211> 4860
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<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

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tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

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gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

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gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

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Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
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Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
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Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350
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Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
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Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
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Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620
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Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
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Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
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Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
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Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895
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Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
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Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
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Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
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Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
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Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
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Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
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Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
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Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
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Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
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Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
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Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
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Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
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Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
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Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
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Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
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Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
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Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
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Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Gly
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Ala Pro  Gly Ser Ser Pro Ser  Ala Ser Thr Gly Thr  Gly Pro Gly
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Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly Pro Gly  Ala Ser Pro
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Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
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Ser Thr  Gly Ser Pro Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly
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Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr Gly Thr  Gly Pro Gly
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Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Thr Pro Gly  Ser Gly Thr
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Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly
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Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
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Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
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Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
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Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
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ggaaatcaag actccttcac acctgtggtg aactctctag acccaccgtt actgactcgc 4680

taccttcgaa ttcaccccca gagttgggtg caccagattg ccctgaggat ggaggttctg 4740

ggctgcgagg cacaggacct ctacggcgcc ggatcacctg gggccgaaac ggccgaacaa 4800

aaactcatct cagaagagga tctgtcacct gcaaccggtc atcatcacca tcaccattga 4860

<210> 707
<211> 1609
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 707
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

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Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400
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```
Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
               405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
               485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
               565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
               645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670
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Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940
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Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1025                1030                1035

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1040                1045                1050

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1055                1060                1065

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1070                1075                1080

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1085                1090                1095

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1100                1105                1110

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1115                1120                1125

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1130                1135                1140

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1145                1150                1155

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1160                1165                1170

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1175                1180                1185

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1190                1195                1200
```

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Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1205                 1210                 1215

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1220                 1225                 1230

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1235                 1240                 1245

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1250                 1255                 1260

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1265                 1270                 1275

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1280                 1285                 1290

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1295                 1300                 1305

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1310                 1315                 1320

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1325                 1330                 1335

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1340                 1345                 1350

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1355                 1360                 1365

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1370                 1375                 1380

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1385                 1390                 1395

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1400                 1405                 1410

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1415                 1420                 1425

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Thr Ser
    1430                 1435                 1440

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala
    1445                 1450                 1455
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Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
1460 1465 1470

Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
1475 1480 1485

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu
1490 1495 1500

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser
1505 1510 1515

Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
1520 1525 1530

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
1535 1540 1545

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser
1550 1555 1560

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro
1565 1570 1575

Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Glu Gln Lys Leu Ile
1580 1585 1590

Ser Glu Glu Asp Leu Ser Pro Ala Thr Gly His His His His His
1595 1600 1605

His

<210> 708
<211> 4830
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 708
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ctcggtgagc tgcctgtgga cgcaagattt cctcctagag tgccaaaatc ttttccattc 120

aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact actttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

agagatctag cttcaggact cattggccct ctcctcatct gctacaaaga atctgtagat 1680

caaagaggaa accagataat gtcagacaag aggaatgtca tcctgttttc tgtatttgat 1740

gagaaccgaa gctggtacct cacagagaat atacaacgct ttctccccaa tccagctgga 1800

gtgcagcttg aggatccaga gttccaagcc tccaacatca tgcacagcat caatggctat 1860

gtttttgata gtttgcagtt gtcagtttgt ttgcatgagg tggcatactg gtacattcta 1920

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cttcagtcag atcaagagga aatcgattat gatgatacca tatcagttga aatgaagaag 2340

gaagattttg acatttatga tgaggatgaa aatcagagcc cccgcagctt tcaaaagaaa 2400

acacgacact attttattgc tgcagtggag aggctctggg attatgggat gagtagctcc 2460

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gatcagaggc aaggagcaga acctagaaaa aactttgtca agcctaatga aaccaaaact 2760

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aattatcgct tccatgcaat caatggctac ataatggata cactacctgg cttagtaatg 3120

gctcaggatc aaaggattcg atggtatctg ctcagcatgg gcagcaatga aaacatccat 3180

tctattcatt tcagtggaca tgtgttcact gtacgaaaaa aagaggagta taaaatggca 3240

ctgtacaatc tctatccagg tgtttttgag acagtggaaa tgttaccatc caaagctgga 3300

atttggcggg tggaatgcct tattggcgag catctacatg ctgggatgag cacacttttt 3360

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ctgttggcac caatgattat tcacggcatc aagacccagg gtgcccgtca gaagttctcc 3600

agcctctaca tctctcagtt tatcatcatg tatagtcttg atgggaagaa gtggcagact 3660

tatcgaggaa attccactgg aaccttaatg gtcttctttg gcaatgtgga ttcatctggg 3720

ataaaacaca atatttttaa ccctccaatt attgctcgat acatccgttt gcacccaact 3780

cattatagca ttcgcagcac tcttcgcatg gagttgatgg gctgtgattt aaatagttgc 3840

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tactttacca atatgtttgc cacctggtct ccttcaaaag ctcgacttca cctccaaggg 3960

aggagtaatg cctggagacc tcaggtgaat aatccaaaag agtggctgca agtggacttc 4020

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atgtatgtga aggagttcct catctccagc agtcaagatg gccatcagtg gactctcttt 4140

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ggctcagaac ccgctacctc aggctccgag acacccggct ctcctgctgg gagtcccact 4560

tccaccgagg aaggaacatc cactgagcct agtgagggct ctgcccctgg aaccagcaca 4620

gagccaagtg agggcagtgc accaggatcc gagccagcaa ccagcgggtc cgagactccc 4680

gggacctctg agtctgccac cccagagagc ggacccggca cttcaaccga gccctccgaa 4740

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gcaaccggtc atcatcacca tcaccattga 4830

<210> 709
<211> 1609
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 709

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu

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            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
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    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
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725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
740 745 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
755 760 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
770 775 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785 790 795 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
805 810 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
820 825 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
835 840 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
850 855 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865 870 875 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
885 890 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
900 905 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
915 920 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
930 935 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945 950 955 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
965 970 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
980 985 990

Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro

995 1000 1005

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1010 1015 1020

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1025 1030 1035

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1040 1045 1050

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1055 1060 1065

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1070 1075 1080

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1085 1090 1095

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1100 1105 1110

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1115 1120 1125

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1130 1135 1140

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1145 1150 1155

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1160 1165 1170

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1175 1180 1185

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1190 1195 1200

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1205 1210 1215

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1220 1225 1230

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1235 1240 1245

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser

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    1250                1255                1260

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1265                1270                1275

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1280                1285                1290

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1295                1300                1305

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1310                1315                1320

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1325                1330                1335

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1340                1345                1350

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1355                1360                1365

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1370                1375                1380

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1385                1390                1395

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1400                1405                1410

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1415                1420                1425

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Pro Gly
    1430                1435                1440

Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly Pro Gly  Ser Ser Pro
    1445                1450                1455

Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Thr Pro Gly  Ser Gly Thr
    1460                1465                1470

Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly
    1475                1480                1485

Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr Gly Thr  Gly Pro Gly
    1490                1495                1500

Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser Pro Gly  Thr Pro Gly
```

1505 1510 1515

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
1520 1525 1530

Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
1535 1540 1545

Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
1550 1555 1560

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
1565 1570 1575

Ser Gly Thr Ala Ser Ser Ser Ala Ser Ser Glu Gln Lys Leu Ile
1580 1585 1590

Ser Glu Glu Asp Leu Ser Pro Ala Thr Gly His His His His His
1595 1600 1605

His

<210> 710
<211> 4830
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 710
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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

agagatctag cttcaggact cattggccct ctcctcatct gctacaaaga atctgtagat 1680

caaagaggaa accagataat gtcagacaag aggaatgtca tcctgttttc tgtatttgat 1740

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agcttctctc aaaacccacc agtcttgaaa cgccatcaac gggaaataac tcgtactact 2280

cttcagtcag atcaagagga aatcgattat gatgatacca tatcagttga aatgaagaag 2340

gaagattttg acatttatga tgaggatgaa aatcagagcc cccgcagctt tcaaaagaaa 2400

acacgacact attttattgc tgcagtggag aggctctggg attatgggat gagtagctcc 2460

ccacatgttc taagaaacag ggctcagagt ggcagtgtcc ctcagttcaa gaaagttgtt 2520

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gcaaccggtc atcatcacca tcaccattga 4830

<210> 711
<211> 1763
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 711

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Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240
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Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510
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Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Pro Gly Ala Ser
            740                 745                 750

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
        755                 760                 765

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
    770                 775                 780
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Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro
785                 790                 795                 800

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
                805                 810                 815

Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
            820                 825                 830

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
        835                 840                 845

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
    850                 855                 860

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
865                 870                 875                 880

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
                885                 890                 895

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
            900                 905                 910

Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
        915                 920                 925

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
    930                 935                 940

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
945                 950                 955                 960

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
                965                 970                 975

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
            980                 985                 990

Pro Gly Thr Ser Ser Thr Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser
        995                 1000                1005

Thr Gly  Thr Gly Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
    1010                1015                1020

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Ala Ser
    1025                1030                1035

Ser Pro  Pro Val Leu Lys Arg  His Gln Arg Glu Ile  Thr Arg Thr
    1040                1045                1050
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Thr Leu  Gln Ser Asp Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile
    1055                1060                1065

Ser Val  Glu Met Lys Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp
    1070                1075                1080

Glu Asn  Gln Ser Pro Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr
    1085                1090                1095

Phe Ile  Ala Ala Val Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser
    1100                1105                1110

Ser Pro  His Val Leu Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro
    1115                1120                1125

Gln Phe  Lys Lys Val Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe
    1130                1135                1140

Thr Gln  Pro Leu Tyr Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu
    1145                1150                1155

Leu Gly  Pro Tyr Ile Arg Ala  Glu Val Glu Asp Asn  Ile Met Val
    1160                1165                1170

Thr Phe  Arg Asn Gln Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser
    1175                1180                1185

Leu Ile  Ser Tyr Glu Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg
    1190                1195                1200

Lys Asn  Phe Val Lys Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys
    1205                1210                1215

Val Gln  His His Met Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys
    1220                1225                1230

Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu Glu Lys  Asp Val His
    1235                1240                1245

Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys His Thr  Asn Thr Leu
    1250                1255                1260

Asn Pro  Ala His Gly Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu
    1265                1270                1275

Phe Phe  Thr Ile Phe Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu
    1280                1285                1290

Asn Met  Glu Arg Asn Cys Arg  Ala Pro Cys Asn Ile  Gln Met Glu
    1295                1300                1305
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Asp Pro  Thr Phe Lys Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly
    1310                1315                1320

Tyr Ile  Met Asp Thr Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln
    1325                1330                1335

Arg Ile  Arg Trp Tyr Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile
    1340                1345                1350

His Ser  Ile His Phe Ser Gly  His Val Phe Thr Val  Arg Lys Lys
    1355                1360                1365

Glu Glu  Tyr Lys Met Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe
    1370                1375                1380

Glu Thr  Val Glu Met Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val
    1385                1390                1395

Glu Cys  Leu Ile Gly Glu His  Leu His Ala Gly Met  Ser Thr Leu
    1400                1405                1410

Phe Leu  Val Tyr Ser Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala
    1415                1420                1425

Ser Gly  His Ile Arg Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr
    1430                1435                1440

Gly Gln  Trp Ala Pro Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser
    1445                1450                1455

Ile Asn  Ala Trp Ser Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val
    1460                1465                1470

Asp Leu  Leu Ala Pro Met Ile  Ile His Gly Ile Lys  Thr Gln Gly
    1475                1480                1485

Ala Arg  Gln Lys Phe Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile
    1490                1495                1500

Met Tyr  Ser Leu Asp Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn
    1505                1510                1515

Ser Thr  Gly Thr Leu Met Val  Phe Phe Gly Asn Val  Asp Ser Ser
    1520                1525                1530

Gly Ile  Lys His Asn Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr
    1535                1540                1545

Ile Arg  Leu His Pro Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg
    1550                1555                1560
```

Met Glu Leu Met Gly Cys Asp Leu Asn Ser Cys Ser Met Pro Leu
1565 1570 1575

Gly Met Glu Ser Lys Ala Ile Ser Asp Ala Gln Ile Thr Ala Ser
1580 1585 1590

Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp Ser Pro Ser Lys Ala
1595 1600 1605

Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg Pro Gln Val
1610 1615 1620

Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln Lys Thr Met
1625 1630 1635

Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser Leu Leu Thr
1640 1645 1650

Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly
1655 1660 1665

His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val Lys Val Phe
1670 1675 1680

Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn Ser Leu Asp
1685 1690 1695

Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro Gln Ser Trp
1700 1705 1710

Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly Cys Glu Ala
1715 1720 1725

Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu Thr Ala Glu
1730 1735 1740

Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala Thr Gly His
1745 1750 1755

His His His His His
1760

<210> 712
<211> 5292
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 712

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<211> 1763
<212> PRT
<213> Artificial Sequence

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

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Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
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Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335
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Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605
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Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Pro Gly Ala Ser
            740                 745                 750

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
        755                 760                 765

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
    770                 775                 780

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro
785                 790                 795                 800

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
                805                 810                 815

Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
            820                 825                 830

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
        835                 840                 845

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
    850                 855                 860

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
865                 870                 875                 880
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Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
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Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
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Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
        915                 920                 925

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
    930                 935                 940

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
945                 950                 955                 960

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
                965                 970                 975

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
            980                 985                 990

Pro Gly Thr Ser Ser Thr Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser
        995                 1000                1005

Thr Gly  Thr Gly Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
    1010                1015                1020

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Ala Ser
    1025                1030                1035

Ser Pro  Pro Val Leu Lys Arg  His Gln Ala Glu Ile  Thr Arg Thr
    1040                1045                1050

Thr Leu  Gln Ser Asp Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile
    1055                1060                1065

Ser Val  Glu Met Lys Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp
    1070                1075                1080

Glu Asn  Gln Ser Pro Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr
    1085                1090                1095

Phe Ile  Ala Ala Val Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser
    1100                1105                1110

Ser Pro  His Val Leu Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro
    1115                1120                1125

Gln Phe  Lys Lys Val Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe
    1130                1135                1140
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Thr Gln  Pro Leu Tyr Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu
    1145                1150                1155

Leu Gly  Pro Tyr Ile Arg Ala  Glu Val Glu Asp Asn  Ile Met Val
    1160                1165                1170

Thr Phe  Arg Asn Gln Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser
    1175                1180                1185

Leu Ile  Ser Tyr Glu Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg
    1190                1195                1200

Lys Asn  Phe Val Lys Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys
    1205                1210                1215

Val Gln  His His Met Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys
    1220                1225                1230

Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu Glu Lys  Asp Val His
    1235                1240                1245

Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys His Thr  Asn Thr Leu
    1250                1255                1260

Asn Pro  Ala His Gly Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu
    1265                1270                1275

Phe Phe  Thr Ile Phe Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu
    1280                1285                1290

Asn Met  Glu Arg Asn Cys Arg  Ala Pro Cys Asn Ile  Gln Met Glu
    1295                1300                1305

Asp Pro  Thr Phe Lys Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly
    1310                1315                1320

Tyr Ile  Met Asp Thr Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln
    1325                1330                1335

Arg Ile  Arg Trp Tyr Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile
    1340                1345                1350

His Ser  Ile His Phe Ser Gly  His Val Phe Thr Val  Arg Lys Lys
    1355                1360                1365

Glu Glu  Tyr Lys Met Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe
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Glu Thr  Val Glu Met Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val
    1385                1390                1395
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Glu Cys  Leu Ile Gly Glu His  Leu His Ala Gly Met  Ser Thr Leu
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Phe Leu  Val Tyr Ser Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala
    1415                1420                1425

Ser Gly  His Ile Arg Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr
    1430                1435                1440

Gly Gln  Trp Ala Pro Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser
    1445                1450                1455

Ile Asn  Ala Trp Ser Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val
    1460                1465                1470

Asp Leu  Leu Ala Pro Met Ile  Ile His Gly Ile Lys  Thr Gln Gly
    1475                1480                1485

Ala Arg  Gln Lys Phe Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile
    1490                1495                1500

Met Tyr  Ser Leu Asp Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn
    1505                1510                1515

Ser Thr  Gly Thr Leu Met Val  Phe Phe Gly Asn Val  Asp Ser Ser
    1520                1525                1530

Gly Ile  Lys His Asn Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr
    1535                1540                1545

Ile Arg  Leu His Pro Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg
    1550                1555                1560

Met Glu  Leu Met Gly Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu
    1565                1570                1575

Gly Met  Glu Ser Lys Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser
    1580                1585                1590

Ser Tyr  Phe Thr Asn Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala
    1595                1600                1605

Arg Leu  His Leu Gln Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val
    1610                1615                1620

Asn Asn  Pro Lys Glu Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met
    1625                1630                1635

Lys Val  Thr Gly Val Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr
    1640                1645                1650
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Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Gln Asp Gly
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His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val Lys Val Phe
1670 1675 1680

Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn Ser Leu Asp
1685 1690 1695

Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro Gln Ser Trp
1700 1705 1710

Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly Cys Glu Ala
1715 1720 1725

Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu Thr Ala Glu
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Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala Thr Gly His
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His His His His His
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

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His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430
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Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700
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Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975
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Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
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Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1025                1030                1035

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1040                1045                1050

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1055                1060                1065

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1070                1075                1080

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1085                1090                1095

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1100                1105                1110

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1115                1120                1125

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1130                1135                1140

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1145                1150                1155

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1160                1165                1170

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1175                1180                1185

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1190                1195                1200

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
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Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1220                1225                1230
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Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
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Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
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Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1265                 1270                 1275

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
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Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1295                 1300                 1305

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
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Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
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Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
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Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
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Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
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Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
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Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
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Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1415                 1420                 1425

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
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Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
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Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
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Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
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Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1520                 1525                 1530

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
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Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
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Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
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Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1595                 1600                 1605

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
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Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
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Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
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Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
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Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Glu Pro Ala
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Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
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Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    1715                 1720                 1725

Pro Ala  Ser Ser Glu Gln Lys  Leu Ile Ser Glu Glu  Asp Leu Ser
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aactctctag acccaccgtt actgactcgc taccttcgaa ttcaccccca gagttgggtg 4260

caccagattg ccctgaggat ggaggttctg ggctgcgagg cacaggacct ctacggcgcg 4320

ccaggtacct cagagtctgc tacccccgag tcagggccag gatcagagcc agccacctcc 4380

gggtctgaga cacccgggac ttccgagagt gccacccctg agtccggacc cgggtccgag 4440

cccgccactt ccggctccga aactcccggc acaagcgaga gcgctacccc agagtcagga 4500

ccaggaacat ctacagagcc ctctgaaggc tccgctccag ggtccccagc cggcagtccc 4560

actagcaccg aggagggaac ctctgaaagc gccacacccg aatcagggcc agggtctgag 4620

cctgctacca gcggcagcga gacaccaggc acctctgagt ccgccacacc agagtccgga 4680

cccggatctc ccgctgggag ccccacctcc actgaggagg gatctcctgc tggctctcca 4740

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ggcagtgaga cacccggcac ttcagagagt gccacccccg agagtggccc aggcactagt 5160

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<210> 717
<211> 1753
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 717
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285
```

```
Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560
```

```
Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830
```

```
Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1025                1030                1035

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1040                1045                1050

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1055                1060                1065

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1070                1075                1080

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1085                1090                1095
```

```
Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1100                1105                1110

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1115                1120                1125

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1130                1135                1140

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1145                1150                1155

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1160                1165                1170

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1175                1180                1185

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1190                1195                1200

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1205                1210                1215

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1220                1225                1230

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1235                1240                1245

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1250                1255                1260

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1265                1270                1275

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1280                1285                1290

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1295                1300                1305

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1310                1315                1320

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1325                1330                1335

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1340                1345                1350
```

```
Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1355                 1360                 1365

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1370                 1375                 1380

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1385                 1390                 1395

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1400                 1405                 1410

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1415                 1420                 1425

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Pro Gly
    1430                 1435                 1440

Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser Pro Gly  Ala Ser Pro
    1445                 1450                 1455

Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Thr Pro Gly  Ser Gly Thr
    1460                 1465                 1470

Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly
    1475                 1480                 1485

Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly
    1490                 1495                 1500

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Thr Pro Gly
    1505                 1510                 1515

Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser Ser Thr  Pro Ser Gly
    1520                 1525                 1530

Ala Thr  Gly Ser Pro Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly
    1535                 1540                 1545

Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr Gly Thr  Gly Pro Gly
    1550                 1555                 1560

Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly Pro Gly  Ala Ser Pro
    1565                 1570                 1575

Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Thr Pro Gly  Ser Gly Thr
    1580                 1585                 1590

Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly
    1595                 1600                 1605
```

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
1610 1615 1620

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
1625 1630 1635

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
1640 1645 1650

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
1655 1660 1665

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
1670 1675 1680

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
1685 1690 1695

Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr
1700 1705 1710

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
1715 1720 1725

Ser Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
1730 1735 1740

Pro Ala Thr Gly His His His His His His
1745 1750

<210> 718
<211> 5262
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

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<210> 719
<211> 2329
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

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Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
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Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415
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Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685
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Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
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```

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Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
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Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1025                1030                1035

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1040                1045                1050

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
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Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
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Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
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Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
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Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1115                1120                1125

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1130                1135                1140

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
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Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
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Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
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Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
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Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
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Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
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Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
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Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
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Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
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Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
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Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
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Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
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Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
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Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
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Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
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Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1400                1405                1410

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1415                1420                1425

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Ser
    1430                1435                1440

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
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Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu
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Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
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Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
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Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Thr  Ser Glu Ser
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Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
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Ser Glu  Thr Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
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Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Thr  Ser Thr Glu
    1565                 1570                 1575

Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Thr Glu  Pro Ser Glu
    1580                 1585                 1590

Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    1595                 1600                 1605

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
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Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Thr  Ser Glu Ser
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Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu
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Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
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Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr
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Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Thr  Ser Thr Glu
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Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
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Glu Ser  Gly Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
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Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
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Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
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Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
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Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
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Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Thr  Ser Thr Glu
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Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Thr Glu  Pro Ser Glu
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Gly Ser  Ala Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
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Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr
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Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Thr  Ser Glu Ser
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Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
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Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
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Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Thr  Ser Thr Glu
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Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1940                1945                1950

Glu Ser  Gly Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    1955                1960                1965

Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser
    1970                1975                1980
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Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
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Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu
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Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
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Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    2045                 2050                 2055

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2060                 2065                 2070

Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    2075                 2080                 2085

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr
    2090                 2095                 2100

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    2105                 2110                 2115

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
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Glu Ser  Gly Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
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Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr
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Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Thr  Ser Glu Ser
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Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2180                 2185                 2190

Glu Ser  Gly Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2195                 2200                 2205

Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Ser
    2210                 2215                 2220

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Ser  Pro Ala Gly
    2225                 2230                 2235
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Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr
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His

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20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

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Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495
```

```
His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765
```

```
Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1025                1030                1035
```

```
Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
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Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1055                1060                1065

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1070                1075                1080

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1085                1090                1095

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1100                1105                1110

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1115                1120                1125

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1130                1135                1140

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1145                1150                1155

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1160                1165                1170

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1175                1180                1185

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1190                1195                1200

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1205                1210                1215

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1220                1225                1230

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1235                1240                1245

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1250                1255                1260

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1265                1270                1275

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1280                1285                1290
```

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Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1310                1315                1320

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1325                1330                1335

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1340                1345                1350

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1355                1360                1365

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1370                1375                1380

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1385                1390                1395

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1400                1405                1410

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1415                1420                1425

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Ala
    1430                1435                1440

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ser  Ser Pro Ser
    1445                1450                1455

Ala Ser  Thr Gly Thr Gly Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1460                1465                1470

Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1475                1480                1485

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser
    1490                1495                1500

Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro Gly Ala  Ser Pro Gly
    1505                1510                1515

Thr Ser  Ser Thr Gly Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala
    1520                1525                1530

Ser Ser  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1535                1540                1545
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Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala
    1550                1555                1560

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1565                1570                1575

Thr Ser  Ser Thr Gly Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala
    1580                1585                1590

Ser Ser  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1595                1600                1605

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Thr
    1610                1615                1620

Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro
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Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
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Gly Thr  Gly Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
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Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser
    1670                1675                1680

Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1685                1690                1695

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1700                1705                1710

Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    1715                1720                1725

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala
    1730                1735                1740

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1745                1750                1755

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
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Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
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Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala
    1790                1795                1800
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Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
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Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1820                 1825                 1830

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1835                 1840                 1845

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ala
    1850                 1855                 1860

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Thr  Pro Gly Ser
    1865                 1870                 1875

Gly Thr  Ala Ser Ser Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1880                 1885                 1890

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1895                 1900                 1905

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser
    1910                 1915                 1920

Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro Gly Ala  Ser Pro Gly
    1925                 1930                 1935

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1940                 1945                 1950

Thr Gly  Ser Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1955                 1960                 1965

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1970                 1975                 1980

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1985                 1990                 1995

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    2000                 2005                 2010

Thr Gly  Ser Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
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Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Thr
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Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro
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Ser Gly  Ala Thr Gly Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala
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Ser Ser  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
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Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser
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Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro Gly Ser  Ser Pro Ser
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Ala Ser  Thr Gly Thr Gly Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    2120                 2125                 2130

Thr Gly  Ser Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    2135                 2140                 2145

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser
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Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro Gly Ser  Ser Pro Ser
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Ala Ser  Thr Gly Thr Gly Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
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Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
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Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser
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Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro Gly Ala  Ser Pro Gly
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Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
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Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    2255                 2260                 2265

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser
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Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
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Thr Ser  Ser Thr Gly Ser Pro  Ala Ser Ser Glu Gln  Lys Leu Ile
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Ser Glu Glu Asp Leu Ser Pro Ala Thr Gly His His His His His
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His

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atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

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acaccgtcgg gagctacggg aagcccagga gcgtcaccgg gaacgtcgtc aacggggtca 4860

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<210> 723
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala
20 25 30

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Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu
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Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
    50                  55                  60

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser
65                  70                  75                  80

Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
                85                  90                  95

Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr
            100                 105                 110

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser
        115                 120                 125

Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr
    130                 135                 140

Glu Glu Gly Ala Ser Ser Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val
145                 150                 155                 160

Glu Leu Ser Trp Asp Tyr Met Gln Ser Asp Leu Gly Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300
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Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575
```

```
Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845
```

```
Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110
```

```
Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365
```

```
Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                1435                1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                1450                1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                1465                1470

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                1480                1485

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1490                1495                1500

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                1510                1515

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1520                1525                1530

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1535                1540                1545

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1550                1555                1560

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1565                1570                1575

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1580                1585                1590

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
    1595                1600                1605

Ser Pro  Ala Thr Gly His His  His His His His His
    1610                1615
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<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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gagcctgcca ctagcggctc cgagactccg ggaacttccg agagcgctac accagaaagc 180

ggacccggaa ccagtaccga acctagcgag ggctctgctc cgggcagccc agccggctct 240

cctacatcca cggaggaggg cacttccgaa tccgccaccc cggagtcagg gccaggatct 300

gaacccgcta cctcaggcag tgagacgcca ggaacgagcg agtccgctac accggagagt 360

gggccaggga gccctgctgg atctcctacg tccactgagg aagggtcacc agcgggctcg 420

cccaccagca ctgaagaagg tgcctcgagc gccaccagaa gatactacct gggtgcagtg 480

gaactgtcat gggactatat gcaaagtgat ctcggtgagc tgcctgtgga cgcaagattt 540

cctcctagag tgccaaaatc ttttccattc aacacctcag tcgtgtacaa aaagactctg 600

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ctgctaggtc ctaccatcca ggctgaggtt tatgatacag tggtcattac acttaagaac 720

atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag 780

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ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

ggcctcattg gagccctact agtatgtaga gaagggagtc tggccaagga aaagacacag 1020

accttgcaca aatttatact acttttgct gtatttgatg aagggaaaag ttggcactca 1080

gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg 1200

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ggctgcgagg cacaggacct ctacggcgcc ggatcacctg gggccgaaac ggccgaacaa 4800

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<210> 725
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 725
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Gly Ala Pro Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
35 40 45

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala
50 55 60

Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu
65 70 75 80

Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr

85 90 95

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser
100 105 110

Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
115 120 125

Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr
130 135 140

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser
145 150 155 160

Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr
165 170 175

Glu Glu Gly Ala Ser Ser Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
260 265 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
275 280 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
290 295 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305 310 315 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
325 330 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
340 345 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln

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        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
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625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
645 650 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
835 840 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
885 890 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln

900 905 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
915 920 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
930 935 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945 950 955 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
965 970 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
980 985 990

Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
995 1000 1005

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1010 1015 1020

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1025 1030 1035

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1040 1045 1050

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1055 1060 1065

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1070 1075 1080

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1085 1090 1095

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1100 1105 1110

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1115 1120 1125

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1130 1135 1140

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1145 1150 1155

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg

1160 1165 1170

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1175 1180 1185

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1190 1195 1200

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1205 1210 1215

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1220 1225 1230

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1235 1240 1245

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1250 1255 1260

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1265 1270 1275

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1280 1285 1290

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1295 1300 1305

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1310 1315 1320

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1325 1330 1335

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1340 1345 1350

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1355 1360 1365

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1370 1375 1380

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1385 1390 1395

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1400 1405 1410

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn

1415 1420 1425

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1430 1435 1440

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1445 1450 1455

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1460 1465 1470

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1475 1480 1485

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1490 1495 1500

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1505 1510 1515

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1520 1525 1530

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1580 1585 1590

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1595 1600 1605

Ser Pro Ala Thr Gly His His His His His His
1610 1615

<210> 726
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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ctcggtgagc tgcctgtgga cgcaagattt cctcctggcg cgccaacatc agagagcgcc 120

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agtgagtctg caactcccga gtccggacct ggctccgagc ctgccactag cggctccgag      240
actccgggaa cttccgagag cgctacacca gaaagcggac ccggaaccag taccgaacct      300
agcgagggct ctgctccggg cagcccagcc ggctctccta catccacgga ggagggcact      360
tccgaatccg ccaccccgga gtcagggcca ggatctgaac ccgctacctc aggcagtgag      420
acgccaggaa cgagcgagtc cgctacaccg gagagtgggc cagggagccc tgctggatct      480
cctacgtcca ctgaggaagg gtcaccagcg ggctcgccca ccagcactga agaaggtgcc      540
tcgagcagag tgccaaaatc ttttccattc aacacctcag tcgtgtacaa aaagactctg      600
tttgtagaat tcacggatca ccttttcaac atcgctaagc caaggccacc ctggatgggt      660
ctgctaggtc ctaccatcca ggctgaggtt tatgatacag tggtcattac acttaagaac      720
atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag      780
ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct      840
ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac      900
ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca      960
ggcctcattg gagccctact agtatgtaga gaagggagtc tggccaagga aaagacacag     1020
accttgcaca aatttatact acttttgct gtatttgatg aagggaaaag ttggcactca     1080
gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa     1140
atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg     1200
aaatcagtct attggcatgt gattggaatg ggcaccactc ctgaagtgca ctcaatattc     1260
ctcgaaggtc acacatttct tgtgaggaac catcgccagg ctagcttgga aatctcgcca     1320
ataactttcc ttactgctca aacactcttg atggaccttg gacagtttct actgttttgt     1380
catatctctt cccaccaaca tgatggcatg gaagcttatg tcaaagtaga cagctgtcca     1440
gaggaacccc aactacgaat gaaaaataat gaagaagcgg aagactatga tgatgatctt     1500
actgattctg aaatggatgt ggtcaggttt gatgatgaca actctccttc ctttatccaa     1560
attcgctcag ttgccaagaa gcatcctaaa acttgggtac attacattgc tgctgaagag     1620
gaggactggg actatgctcc cttagtcctc gcccccgatg acagaagtta taaaagtcaa     1680
tatttgaaca atggccctca gcggattggt aggaagtaca aaaaagtccg atttatggca     1740
tacacagatg aaacctttaa gactcgtgaa gctattcagc atgaatcagg aatcttggga     1800
cctttacttt atggggaagt tggagacaca ctgttgatta tatttaagaa tcaagcaagc     1860
agaccatata acatctaccc tcacggaatc actgatgtcc gtcctttgta ttcaaggaga     1920
ttaccaaaag gtgtaaaaca tttgaaggat tttccaattc tgccaggaga aatattcaaa     1980
tataaatgga cagtgactgt agaagatggg ccaactaaat cagatcctcg gtgcctgacc     2040
cgctattact ctagtttcgt taatatggag agagatctag cttcaggact cattggccct     2100
ctcctcatct gctacaaaga atctgtagat caaagaggaa accagataat gtcagacaag     2160
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atacaacgct ttctccccaa tccagctgga gtgcagcttg aggatccaga gttccaagcc 2280

tccaacatca tgcacagcat caatggctat gtttttgata gtttgcagtt gtcagtttgt 2340

ttgcatgagg tggcatactg gtacattcta agcattggag cacagactga cttcctttct 2400

gtcttcttct ctggatatac cttcaaacac aaaatggtct atgaagacac actcacccta 2460

ttcccattct caggagaaac tgtcttcatg tcgatggaaa acccaggtct atggattctg 2520

gggtgccaca actcagactt tcggaacaga ggcatgaccg ccttactgaa ggtttctagt 2580

tgtgacaaga acactggtga ttattacgag gacagttatg aagatatttc agcatacttg 2640

ctgagtaaaa acaatgccat tgaaccaaga agcttctctc aaaacccacc agtcttgaaa 2700

cgccatcaac gggaaataac tcgtactact cttcagtcag atcaagagga aatcgattat 2760

gatgatacca tatcagttga aatgaagaag gaagattttg acatttatga tgaggatgaa 2820

aatcagagcc cccgcagctt tcaaaagaaa acacgacact attttattgc tgcagtggag 2880

aggctctggg attatgggat gagtagctcc ccacatgttc taagaaacag ggctcagagt 2940

ggcagtgtcc ctcagttcaa gaaagttgtt ttccaggaat ttactgatgg ctcctttact 3000

cagcccttat accgtggaga actaaatgaa catttgggac tcctggggcc atatataaga 3060

gcagaagttg aagataatat catggtaact ttcagaaatc aggcctctcg tccctattcc 3120

ttctattcta gccttatttc ttatgaggaa gatcagaggc aaggagcaga acctagaaaa 3180

aactttgtca agcctaatga aaccaaaact tacttttgga aagtgcaaca tcatatggca 3240

cccactaaag atgagtttga ctgcaaagcc tgggcttatt tctctgatgt tgacctggaa 3300

aaagatgtgc actcaggcct gattggaccc cttctggtct gccacactaa cacactgaac 3360

cctgctcatg ggagacaagt gacagtacag gaatttgctc tgtttttcac catctttgat 3420

gagaccaaaa gctggtactt cactgaaaat atggaaagaa actgcagggc tccctgcaat 3480

atccagatgg aagatcccac ttttaaagag aattatcgct tccatgcaat caatggctac 3540

ataatggata cactacctgg cttagtaatg gctcaggatc aaaggattcg atggtatctg 3600

ctcagcatgg gcagcaatga aaacatccat tctattcatt tcagtggaca tgtgttcact 3660

gtacgaaaaa aagaggagta taaaatggca ctgtacaatc tctatccagg tgtttttgag 3720

acagtggaaa tgttaccatc caaagctgga atttggcggg tggaatgcct tattggcgag 3780

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aagacccagg gtgcccgtca gaagttctcc agcctctaca tctctcagtt tatcatcatg 4080

tatagtcttg atgggaagaa gtggcagact tatcgaggaa attccactgg aaccttaatg 4140

gtcttctttg gcaatgtgga ttcatctggg ataaaacaca atatttttaa ccctccaatt 4200

attgctcgat acatccgttt gcacccaact cattatagca ttcgcagcac tcttcgcatg 4260

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ggaaatcaag actccttcac acctgtggtg aactctctag acccaccgtt actgactcgc 4680

taccttcgaa ttcaccccca gagttgggtg caccagattg ccctgaggat ggaggttctg 4740

ggctgcgagg cacaggacct ctacggcgcc ggatcacctg gggccgaaac ggccgaacaa 4800

aaactcatct cagaagagga tctgtcacct gcaaccggtc atcatcacca tcaccattga 4860

<210> 727
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 727
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Gly Ala Pro Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
65 70 75 80

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
85 90 95

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
100 105 110

Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
115 120 125

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
130 135 140

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Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
145                 150                 155                 160

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
                165                 170                 175

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly
            180                 185                 190

Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser
        195                 200                 205

Thr Glu Glu Gly Ala Ser Ser Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415
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His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685
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Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960
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Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215
```

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Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
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Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                1435                1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                1450                1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                1465                1470
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Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1475 1480 1485

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1490 1495 1500

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1505 1510 1515

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1520 1525 1530

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1580 1585 1590

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1595 1600 1605

Ser Pro Ala Thr Gly His His His His His His
1610 1615

<210> 728
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggggcgc gccaacatca gagagcgcca cccctgaaag tggtcccggg 240

agcgagccag ccacatctgg gtcggaaacg ccaggcacaa gtgagtctgc aactcccgag 300

tccggacctg gctccgagcc tgccactagc ggctccgaga ctccgggaac ttccgagagc 360

gctacaccag aaagcggacc cggaaccagt accgaaccta gcgagggctc tgctccgggc 420

agcccagccg gctctcctac atccacggag gagggcactt ccgaatccgc caccccggag 480

tcagggccag gatctgaacc cgctacctca ggcagtgaga cgccaggaac gagcgagtcc 540

gctacaccgg agagtgggcc agggagccct gctggatctc ctacgtccac tgaggaaggg 600

tcaccagcgg gctcgcccac cagcactgaa gaaggtgcct cgagcccacc ctggatgggt 660

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ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct 840

ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

ggcctcattg gagccctact agtatgtaga gaagggagtc tggccaagga aaagacacag 1020

accttgcaca aatttatact acttttgct gtatttgatg aagggaaaag ttggcactca 1080

gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg 1200

aaatcagtct attggcatgt gattggaatg ggcaccactc ctgaagtgca ctcaatattc 1260

ctcgaaggtc acacatttct tgtgaggaac catcgccagg ctagcttgga aatctcgcca 1320

ataactttcc ttactgctca aacactcttg atggaccttg gacagtttct actgttttgt 1380

catatctctt cccaccaaca tgatggcatg gaagcttatg tcaaagtaga cagctgtcca 1440

gaggaacccc aactacgaat gaaaaataat gaagaagcgg aagactatga tgatgatctt 1500

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attcgctcag ttgccaagaa gcatcctaaa acttgggtac attacattgc tgctgaagag 1620

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cctttacttt atggggaagt tggagacaca ctgttgatta tatttaagaa tcaagcaagc 1860

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aatcagagcc cccgcagctt tcaaaagaaa acacgacact attttattgc tgcagtggag 2880

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cctgctcatg ggagacaagt gacagtacag gaatttgctc tgtttttcac catctttgat 3420

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ggctgcgagg cacaggacct ctacggcgcc ggatcacctg gggccgaaac ggccgaacaa 4800

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<210> 729
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Gly Ala Pro Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
85 90 95

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
100 105 110

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
115 120 125

Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
130 135 140

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
145 150 155 160

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
165 170 175

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
180 185 190

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Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly
        195                 200                 205

Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser
    210                 215                 220

Thr Glu Glu Gly Ala Ser Ser Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460
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His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735
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Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005
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Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260
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```
Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                1435                1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                1450                1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                1465                1470

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                1480                1485

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1490                1495                1500

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                1510                1515
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Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1520 1525 1530

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1580 1585 1590

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1595 1600 1605

Ser Pro Ala Thr Gly His His His His His His
1610 1615

<210> 730
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

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tctcctacat ccacggagga gggcacttcc gaatccgcca ccccggagtc agggccagga 540

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ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

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gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg 1200

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catatctctt cccaccaaca tgatggcatg gaagcttatg tcaaagtaga cagctgtcca 1440

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<212> PRT
<213> Artificial Sequence

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gly Ala Pro Gly Ser Ser Pro Ser Ala
        115                 120                 125

Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr
    130                 135                 140

Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
145                 150                 155                 160

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser
                165                 170                 175

Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr
            180                 185                 190

Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser
        195                 200                 205

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly
    210                 215                 220

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
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```
Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala
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Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Gln Arg Glu
           260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
       275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
   290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
               325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
           340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
       355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
   370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
               405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
           420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
       435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
   450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
               485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
           500                 505                 510
```

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Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
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Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780
```

```
Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
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Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
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Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
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Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
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Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
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Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
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Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
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Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
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```

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Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
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Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
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Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
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Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
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Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
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Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
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Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
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Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
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Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
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Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
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Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
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Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                1510                1515

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1520                1525                1530

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1535                1540                1545

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1550                1555                1560
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Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
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Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
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actcagggag taaaatctct gcttaccagc atgtatgtga aggagttcct catctccagc 4560

agtcaagatg gccatcagtg gactctcttt tttcagaatg gcaaagtaaa ggtttttcag 4620

ggaaatcaag actccttcac acctgtggtg aactctctag acccaccgtt actgactcgc 4680

taccttcgaa ttcaccccca gagttgggtg caccagattg ccctgaggat ggaggttctg 4740

ggctgcgagg cacaggacct ctacggcgcc ggatcacctg gggccgaaac ggccgaacaa 4800

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<210> 733
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 733
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro

20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Gly Ala Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr
210 215 220

Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala
225 230 235 240

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
245 250 255

Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
260 265 270

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala
275 280 285

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala

290 295 300

Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
305 310 315 320

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser
325 330 335

Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
340 345 350

Ser Ser Thr Gly Ser Pro Ala Ser Ser Thr Lys Asn Ser Leu Met Gln
355 360 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
370 375 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385 390 395 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
405 410 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
420 425 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
435 440 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
450 455 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465 470 475 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
485 490 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
500 505 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
515 520 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
530 535 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545 550 555 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val

565 570 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
580 585 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
595 600 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
610 615 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
645 650 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg

```
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
```

1100 1105 1110

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1115 1120 1125

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1130 1135 1140

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1145 1150 1155

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1160 1165 1170

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1175 1180 1185

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1190 1195 1200

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1205 1210 1215

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1220 1225 1230

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1235 1240 1245

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1250 1255 1260

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1265 1270 1275

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1280 1285 1290

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1295 1300 1305

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1310 1315 1320

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1325 1330 1335

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1340 1345 1350

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp

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    1355                1360                1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                1435                1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                1450                1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                1465                1470

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                1480                1485

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1490                1495                1500

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                1510                1515

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1520                1525                1530

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1535                1540                1545

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1550                1555                1560

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1565                1570                1575

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1580                1585                1590

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
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Ser Pro  Ala Thr Gly His His  His His His His
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1610 1615

<210> 734
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaggcgcgc caggctccag cccctccgcg 660

agcacgggaa ccggaccagg ttcgtcaccc tcagcatcaa cggggacggg accgggggcg 720

tcaccaggaa cgtcctccac cggctcgccg ggtgcatcac ccggaacgtc atcgaccgga 780

tcgccaggga gctcgacgcc atcaggcgca acaggatcac ctggctcaag ccctagcgcg 840

tcaaccggca cgggtccggg tgcctcccct ggcacgtcca gcaccggatc acccggatcg 900

agcccatccg cctcaaccgg aaccggaccc ggtacaccag ggtcgggaac agcctcctcg 960

tcaccaggct cctcaacccc ctcgggagcc acgggttcgc ccggttcgtc aacgccttcc 1020

ggagcaactg gtagccccgg agcatcgcca ggaacttcga gcacggggtc gcccgcctcg 1080

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atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg 1200

aaatcagtct attggcatgt gattggaatg ggcaccactc ctgaagtgca ctcaatattc 1260

ctcgaaggtc acacatttct tgtgaggaac catcgccagg ctagcttgga aatctcgcca 1320

ataactttcc ttactgctca aacactcttg atggaccttg gacagtttct actgttttgt 1380

catatctctt cccaccaaca tgatggcatg gaagcttatg tcaaagtaga cagctgtcca 1440

gaggaacccc aactacgaat gaaaaataat gaagaagcgg aagactatga tgatgatctt 1500

actgattctg aaatggatgt ggtcaggttt gatgatgaca actctccttc ctttatccaa 1560

attcgctcag ttgccaagaa gcatcctaaa acttgggtac attacattgc tgctgaagag 1620

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agtcaagatg gccatcagtg gactctcttt tttcagaatg gcaaagtaaa ggtttttcag 4620

ggaaatcaag actccttcac acctgtggtg aactctctag acccaccgtt actgactcgc 4680

taccttcgaa ttcaccccca gagttgggtg caccagattg ccctgaggat ggaggttctg 4740

ggctgcgagg cacaggacct ctacggcgcc ggatcacctg gggccgaaac ggccgaacaa 4800

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<210> 735
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

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Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Gly Ala Pro Gly
    210                 215                 220

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser
225                 230                 235                 240

Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
                245                 250                 255

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
            260                 265                 270

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser
        275                 280                 285

Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
    290                 295                 300

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
305                 310                 315                 320

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
                325                 330                 335

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
            340                 345                 350
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Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala
        355                 360                 365

Ser Ser Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620
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Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895
```

```
Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155
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Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                 1165                 1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                 1180                 1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                 1195                 1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                 1210                 1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                 1225                 1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                 1240                 1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                 1255                 1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                 1270                 1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                 1285                 1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                 1300                 1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                 1315                 1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                 1330                 1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                 1345                 1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                 1360                 1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                 1375                 1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                 1390                 1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                 1405                 1410
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```
Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                 1420                 1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                 1435                 1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                 1450                 1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                 1465                 1470

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                 1480                 1485

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1490                 1495                 1500

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                 1510                 1515

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1520                 1525                 1530

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1535                 1540                 1545

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1550                 1555                 1560

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1565                 1570                 1575

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1580                 1585                 1590

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
    1595                 1600                 1605

Ser Pro  Ala Thr Gly His His  His His His His His
    1610                 1615
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<210> 736
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 736

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

ggcgcgccag gctccagccc ctccgcgagc acgggaaccg gaccaggttc gtcaccctca 720

gcatcaacgg ggacgggacc gggggcgtca ccaggaacgt cctccaccgg ctcgccgggt 780

gcatcacccg gaacgtcatc gaccggatcg ccagggagct cgacgccatc aggcgcaaca 840

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acttcgagca cggggtcgcc cgcctcgagc gatgctgcat ctgctcgggc ctggcctaaa 1140

atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg 1200

aaatcagtct attggcatgt gattggaatg ggcaccactc ctgaagtgca ctcaatattc 1260

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ataactttcc ttactgctca aacactcttg atggaccttg gacagtttct actgttttgt 1380

catatctctt cccaccaaca tgatggcatg gaagcttatg tcaaagtaga cagctgtcca 1440

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attcgctcag ttgccaagaa gcatcctaaa acttgggtac attacattgc tgctgaagag 1620

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cctttacttt atggggaagt tggagacaca ctgttgatta tatttaagaa tcaagcaagc 1860

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tataaatgga cagtgactgt agaagatggg ccaactaaat cagatcctcg gtgcctgacc 2040

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cgccatcaac gggaaataac tcgtactact cttcagtcag atcaagagga aatcgattat 2760

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cagcccttat accgtggaga actaaatgaa catttgggac tcctggggcc atatataaga 3060

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ggctgcgagg cacaggacct ctacggcgcc ggatcacctg gggccgaaac ggccgaacaa 4800

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<210> 737
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 737
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

```
Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Gly Ala Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
225                 230                 235                 240

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly
                245                 250                 255

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
            260                 265                 270

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
        275                 280                 285

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly
    290                 295                 300

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
305                 310                 315                 320

Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
                325                 330                 335

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
            340                 345                 350

Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
        355                 360                 365

Gly Ser Pro Ala Ser Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400
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Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670
```

```
Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940
```

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Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200
```

```
Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                 1210                 1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                 1225                 1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                 1240                 1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                 1255                 1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                 1270                 1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                 1285                 1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                 1300                 1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                 1315                 1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                 1330                 1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                 1345                 1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                 1360                 1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                 1375                 1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                 1390                 1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                 1405                 1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                 1420                 1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                 1435                 1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                 1450                 1455
```

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1460 1465 1470

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1475 1480 1485

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1490 1495 1500

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1505 1510 1515

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1520 1525 1530

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1580 1585 1590

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
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Ser Pro Ala Thr Gly His His His His His His
1610 1615

<210> 738
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

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Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
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His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Gly Ala Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
            340                 345                 350

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly
        355                 360                 365

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
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Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
385                 390                 395                 400

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly
                405                 410                 415

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
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Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
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Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
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Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
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Gly Ser Pro Ala Ser Ser Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
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Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
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Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
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Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
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Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
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Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
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Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
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Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
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Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
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Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
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Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
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Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
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Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
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Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
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Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
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Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
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Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
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Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
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Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
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Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
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Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
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Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
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Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
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Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
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Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
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Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
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Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
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Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
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Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
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Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
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Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
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Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
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Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
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Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
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Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
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Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
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Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
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Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
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Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
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Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
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aactttgtca agcctaatga aaccaaaact tacttttgga aagtgcaaca tcatatggca 3240

cccactaaag atgagtttga ctgcaaagcc tgggcttatt tctctgatgt tgacctggaa 3300

aaagatgtgc actcaggcct gattggaccc cttctggtct gccacactaa cacactgaac 3360

cctgctcatg ggagacaagt gacagtacag gaatttgctc tgtttttcac catctttgat 3420

gagaccaaaa gctggtactt cactgaaaat atggaaagaa actgcagggc tccctgcaat 3480

atccagatgg aagatcccac ttttaaagag aattatcgct tccatgcaat caatggctac 3540

ataatggata cactacctgg cttagtaatg gctcaggatc aaaggattcg atggtatctg 3600

ctcagcatgg gcagcaatga aaacatccat tctattcatt tcagtggaca tgtgttcact 3660

gtacgaaaaa aagaggagta taaaatggca ctgtacaatc tctatccagg tgtttttgag 3720

acagtggaaa tgttaccatc caaagctgga atttggcggg tggaatgcct tattggcgag 3780

catctacatg ctgggatgag cacacttttt ctggtgtaca gcaataagtg tcagactccc 3840

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ggaaatcaag actccttcac acctgtggtg aactctctag acccaccgtt actgactcgc 4680

taccttcgaa ttcaccccca gagttgggtg caccagattg ccctgaggat ggaggttctg 4740

ggctgcgagg cacaggacct ctacggcgcc ggatcacctg gggccgaaac ggccgaacaa 4800

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<210> 741
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 741
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg

225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Gly Ala Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr
340 345 350

Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala
355 360 365

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
370 375 380

Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
385 390 395 400

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala
405 410 415

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala
420 425 430

Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
435 440 445

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser
450 455 460

Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
465 470 475 480

Ser Ser Thr Gly Ser Pro Ala Ser Ser Asn Glu Glu Ala Glu Asp Tyr
485 490 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp

500 505 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
515 520 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
530 535 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545 550 555 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
565 570 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
580 585 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
595 600 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
610 615 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
645 650 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val

770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
835 840 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
885 890 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
900 905 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
915 920 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
930 935 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945 950 955 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
965 970 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
980 985 990

Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
995 1000 1005

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1010 1015 1020

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1025 1030 1035

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg

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    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
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1295 1300 1305

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1310 1315 1320

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1325 1330 1335

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1340 1345 1350

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1355 1360 1365

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1370 1375 1380

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1385 1390 1395

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1400 1405 1410

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1415 1420 1425

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1430 1435 1440

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1445 1450 1455

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1460 1465 1470

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1475 1480 1485

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1490 1495 1500

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1505 1510 1515

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1520 1525 1530

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg

1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1580 1585 1590

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1595 1600 1605

Ser Pro Ala Thr Gly His His His His His His
1610 1615

<210> 742
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

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cctttacttt atggggaagt tggagacaca ctgttgatta tatttaagaa tcaagcaagc 1860

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ctcagcatgg gcagcaatga aaacatccat tctattcatt tcagtggaca tgtgttcact 3660

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ggctgcgagg cacaggacct ctacggcgcc ggatcacctg gggccgaaac ggccgaacaa 4800

aaactcatct cagaagagga tctgtcacct gcaaccggtc atcatcacca tcaccattga 4860

<210> 743
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285
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Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Gly Ala Pro Thr Ser Thr
    370                 375                 380

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
385                 390                 395                 400

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
                405                 410                 415

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
            420                 425                 430

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
        435                 440                 445

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
    450                 455                 460

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr
465                 470                 475                 480

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
                485                 490                 495

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
            500                 505                 510

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560
```

```
Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830
```

```
Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095
```

```
Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350
```

```
Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                1435                1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                1450                1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                1465                1470

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                1480                1485

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1490                1495                1500

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                1510                1515

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1520                1525                1530

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1535                1540                1545

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1550                1555                1560

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1565                1570                1575

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1580                1585                1590

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
    1595                1600                1605
```

Ser Pro Ala Thr Gly His His His His His His
1610 1615

<210> 744
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatggcgcg 1140

ccaaccagta cggagccgtc cgaggggagc gcaccaggaa gcccggctgg gagcccgact 1200

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gaatcagggc caggatctga gcccgcgact tcgggctccg agacgcctgg gacatccacc 1440

gagccctccg aaggatcagc cccaggcacc agcacggagc cctctgaggg aagcgcacct 1500

ggtaccagcg aaagcgcaac tcccgaatca ggtcccggta cgagcgagtc ggcgaccccg 1560

gagagcgggc caggtgcctc gagccctaaa acttgggtac attacattgc tgctgaagag 1620

gaggactggg actatgctcc cttagtcctc gcccccgatg acagaagtta taaaagtcaa 1680

tatttgaaca atggccctca gcggattggt aggaagtaca aaaaagtccg atttatggca 1740

tacacagatg aaacctttaa gactcgtgaa gctattcagc atgaatcagg aatcttggga 1800

cctttacttt atggggaagt tggagacaca ctgttgatta tatttaagaa tcaagcaagc 1860

agaccatata acatctaccc tcacggaatc actgatgtcc gtcctttgta ttcaaggaga 1920

ttaccaaaag gtgtaaaaca tttgaaggat tttccaattc tgccaggaga aatattcaaa 1980

tataaatgga cagtgactgt agaagatggg ccaactaaat cagatcctcg gtgcctgacc 2040

cgctattact ctagtttcgt taatatggag agagatctag cttcaggact cattggccct 2100

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aaagatgtgc actcaggcct gattggaccc cttctggtct gccacactaa cacactgaac 3360

cctgctcatg ggagacaagt gacagtacag gaatttgctc tgtttttcac catctttgat 3420

gagaccaaaa gctggtactt cactgaaaat atggaaagaa actgcagggc tccctgcaat 3480

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ataatggata cactacctgg cttagtaatg gctcaggatc aaaggattcg atggtatctg 3600

ctcagcatgg gcagcaatga aaacatccat tctattcatt tcagtggaca tgtgttcact 3660

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ggctgcgagg cacaggacct ctacggcgcc ggatcacctg gggccgaaac ggccgaacaa 4800

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<210> 745
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

```
Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335
```

```
Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Gly
385                 390                 395                 400

Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro
                405                 410                 415

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
            420                 425                 430

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
        435                 440                 445

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
    450                 455                 460

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
465                 470                 475                 480

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
                485                 490                 495

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
            500                 505                 510

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
        515                 520                 525

Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    530                 535                 540

Pro Gly Ala Ser Ser Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605
```

```
Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880
```

```
Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140
```

```
Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395
```

```
Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                1435                1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                1450                1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                1465                1470

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                1480                1485

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1490                1495                1500

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                1510                1515

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1520                1525                1530

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1535                1540                1545

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1550                1555                1560

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1565                1570                1575

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1580                1585                1590

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
    1595                1600                1605

Ser Pro  Ala Thr Gly His His  His His His His His
    1610                1615
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<210> 746
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic

polynucleotide

<400> 746

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atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

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gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

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<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

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Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380
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Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gly Ala Pro Thr Ser Thr Glu
                405                 410                 415

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
            420                 425                 430

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
        435                 440                 445

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
    450                 455                 460

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
465                 470                 475                 480

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
                485                 490                 495

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
            500                 505                 510

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
        515                 520                 525

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
    530                 535                 540

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655
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Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925
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Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185
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Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                1435                1440
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Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1445 1450 1455

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1460 1465 1470

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1475 1480 1485

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1490 1495 1500

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1505 1510 1515

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1520 1525 1530

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1580 1585 1590

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
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Ser Pro Ala Thr Gly His His His His His His
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<210> 748
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

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ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

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gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

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cctttacttt atggggaagt tggagacaca ctgttgatta tatttaagaa tcaagcaagc 1860

agaccatata acatctaccc tcacggaatc actgatgtcc gtcctttgta ttcaaggaga 1920

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<211> 1619
<212> PRT
<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic polypeptide

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20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu

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Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gly Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
            420                 425                 430

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro
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435 440 445

Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
450 455 460

Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
465 470 475 480

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala
485 490 495

Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu
500 505 510

Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
515 520 525

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala
530 535 540

Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
545 550 555 560

Gly Pro Gly Ala Ser Ser Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
565 570 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
580 585 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
595 600 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
610 615 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
645 650 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys

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705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
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980 985 990

Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
995 1000 1005

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1010 1015 1020

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1025 1030 1035

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1040 1045 1050

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1055 1060 1065

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1070 1075 1080

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1085 1090 1095

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1100 1105 1110

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1115 1120 1125

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1130 1135 1140

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1145 1150 1155

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1160 1165 1170

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1175 1180 1185

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1190 1195 1200

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1205 1210 1215

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1220 1225 1230

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys

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Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
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Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
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Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
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Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                1435                1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                1450                1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                1465                1470

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                1480                1485

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
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1490 1495 1500

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1505 1510 1515

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1520 1525 1530

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1580 1585 1590

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
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Ser Pro Ala Thr Gly His His His His His His
1610 1615

<210> 750
<211> 4860
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<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

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<210> 751
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

```
Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Gly Ala Pro Thr Ser Thr Glu Pro Ser
                485                 490                 495
```

```
Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
            500                 505                 510

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
        515                 520                 525

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
    530                 535                 540

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
545                 550                 555                 560

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
                565                 570                 575

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser
            580                 585                 590

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
        595                 600                 605

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
    610                 615                 620

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765
```

```
Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035
```

```
Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290
```

```
Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                1435                1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                1450                1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                1465                1470

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                1480                1485

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1490                1495                1500

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                1510                1515

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1520                1525                1530

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1535                1540                1545
```

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1580 1585 1590

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1595 1600 1605

Ser Pro Ala Thr Gly His His His His His His
1610 1615

<210> 752
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

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gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta tggcgcgcca accagtacgg agccgtccga ggggagcgca 1500

ccaggaagcc cggctgggag cccgacttct accgaagagg gtacatctac cgaaccaagt 1560

gaaggttcag caccaggcac ctcaacagaa ccctctgagg gctcggcgcc tggtacaagt 1620

gagtccgcca ccccagaatc cgggcctggg acaagcacag aaccttcgga agggagtgcc 1680

cctggaacat ccgaatcggc aaccccagaa tcagggccag gatctgagcc cgcgacttcg 1740

ggctccgaga cgcctgggac atccaccgag ccctccgaag gatcagcccc aggcaccagc 1800

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cccggtacga gcgagtcggc gaccccggag agcgggccag gtgcctcgag ctcaaggaga 1920

ttaccaaaag gtgtaaaaca tttgaaggat tttccaattc tgccaggaga aatattcaaa 1980

tataaatgga cagtgactgt agaagatggg ccaactaaat cagatcctcg gtgcctgacc 2040

cgctattact ctagtttcgt taatatggag agagatctag cttcaggact cattggccct 2100

ctcctcatct gctacaaaga atctgtagat caaagaggaa accagataat gtcagacaag 2160

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tgtgacaaga acactggtga ttattacgag gacagttatg aagatatttc agcatacttg 2640

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cccactaaag atgagtttga ctgcaaagcc tgggcttatt tctctgatgt tgacctggaa 3300

aaagatgtgc actcaggcct gattggaccc cttctggtct gccacactaa cacactgaac 3360

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gagaccaaaa gctggtactt cactgaaaat atggaaagaa actgcagggc tccctgcaat 3480

atccagatgg aagatcccac ttttaaagag aattatcgct tccatgcaat caatggctac 3540

ataatggata cactacctgg cttagtaatg gctcaggatc aaaggattcg atggtatctg 3600

ctcagcatgg gcagcaatga aaacatccat tctattcatt tcagtggaca tgtgttcact 3660

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<210> 753
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 753

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270
```

```
Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Gly Ala
                485                 490                 495

Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
            500                 505                 510

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
        515                 520                 525

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    530                 535                 540
```

```
Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
545                 550                 555                 560

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
                565                 570                 575

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
            580                 585                 590

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
        595                 600                 605

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
    610                 615                 620

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
625                 630                 635                 640

Gly Ala Ser Ser Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815
```

```
Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080
```

```
Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335
```

```
Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                1435                1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                1450                1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                1465                1470

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                1480                1485

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1490                1495                1500

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                1510                1515

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1520                1525                1530

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1535                1540                1545

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1550                1555                1560

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1565                1570                1575

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1580                1585                1590
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Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
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Ser Pro Ala Thr Gly His His His His His His
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<210> 754
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

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catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

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aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

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gagccgtccg aggggagcgc accaggaagc ccggctggga gcccgacttc taccgaagag 1560

ggtacatcta ccgaaccaag tgaaggttca gcaccaggca cctcaacaga accctctgag 1620

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agcgcaactc ccgaatcagg tcccggtacg agcgagtcgg cgaccccgga gagcgggcca 1920

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tataaatgga cagtgactgt agaagatggg ccaactaaat cagatcctcg gtgcctgacc 2040

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ggctgcgagg cacaggacct ctacggcgcc ggatcacctg gggccgaaac ggccgaacaa 4800

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<210> 755
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

```
Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320
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Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Gly Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser
            500                 505                 510

Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr
        515                 520                 525

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro
    530                 535                 540

Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
545                 550                 555                 560

Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
                565                 570                 575

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr
            580                 585                 590
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Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
        595                 600                 605

Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
    610                 615                 620

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala
625                 630                 635                 640

Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860
```

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
885 890 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
900 905 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
915 920 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
930 935 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945 950 955 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
965 970 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
980 985 990

Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
995 1000 1005

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1010 1015 1020

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1025 1030 1035

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1040 1045 1050

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1055 1060 1065

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1070 1075 1080

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1085 1090 1095

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1100 1105 1110

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1115 1120 1125

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Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380
```

```
Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                1435                1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                1450                1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                1465                1470

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                1480                1485

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1490                1495                1500

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                1510                1515

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1520                1525                1530

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1535                1540                1545

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1550                1555                1560

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1565                1570                1575

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1580                1585                1590

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
    1595                1600                1605

Ser Pro  Ala Thr Gly His His  His His His His His
    1610                1615
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<212> DNA
<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic
polynucleotide

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gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala

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Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
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Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
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Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
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Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
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Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Gly Ala Pro Gly Ala
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610 615 620

Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr
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Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser

645 650 655

Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala
660 665 670

Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
675 680 685

Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser
690 695 700

Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly
705 710 715 720

Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
725 730 735

Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser
740 745 750

Ser Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
835 840 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
885 890 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
900 905 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met

915 920 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
930 935 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945 950 955 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
965 970 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
980 985 990

Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
995 1000 1005

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1010 1015 1020

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1025 1030 1035

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
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Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
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Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1070 1075 1080

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
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Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1100 1105 1110

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1115 1120 1125

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1130 1135 1140

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1145 1150 1155

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1160 1165 1170

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu

1175 1180 1185

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1190 1195 1200

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
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Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1220 1225 1230

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1235 1240 1245

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1250 1255 1260

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
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Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1280 1285 1290

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1295 1300 1305

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1310 1315 1320

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1325 1330 1335

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1340 1345 1350

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1355 1360 1365

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
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Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1385 1390 1395

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
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Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1415 1420 1425

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp

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Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
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Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
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Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
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Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
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Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
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atatcagatg cacagattac tgcttcatcc tactttacca atatgtttgc cacctggtct 4380

ccttcaaaag ctcgacttca cctccaaggg aggagtaatg cctggagacc tcaggtgaat 4440

aatccaaaag agtggctgca agtggacttc cagaagacaa tgaaagtcac aggagtaact 4500

actcagggag taaaatctct gcttaccagc atgtatgtga aggagttcct catctccagc 4560

agtcaagatg gccatcagtg gactctcttt tttcagaatg gcaaagtaaa ggtttttcag 4620

ggaaatcaag actccttcac acctgtggtg aactctctag acccaccgtt actgactcgc 4680

taccttcgaa ttcaccccca gagttgggtg caccagattg ccctgaggat ggaggttctg 4740

ggctgcgagg cacaggacct ctacggcgcc ggatcacctg gggccgaaac ggccgaacaa 4800

aaactcatct cagaagagga tctgtcacct gcaaccggtc atcatcacca tcaccattga 4860

<210> 759
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 759
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

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His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430
```

```
Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700
```

```
Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Gly Ala Pro Gly Thr Pro
        755                 760                 765

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser
    770                 775                 780

Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
785                 790                 795                 800

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
                805                 810                 815

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr
            820                 825                 830

Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
        835                 840                 845

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
    850                 855                 860

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly
865                 870                 875                 880

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
                885                 890                 895

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975
```

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Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230
```

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Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                 1240                 1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                 1255                 1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                 1270                 1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                 1285                 1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                 1300                 1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                 1315                 1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                 1330                 1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                 1345                 1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                 1360                 1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                 1375                 1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                 1390                 1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                 1405                 1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                 1420                 1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                 1435                 1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                 1450                 1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                 1465                 1470

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                 1480                 1485
```

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1490 1495 1500

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1505 1510 1515

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1520 1525 1530

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1580 1585 1590

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1595 1600 1605

Ser Pro Ala Thr Gly His His His His His His
1610 1615

<210> 760
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 760
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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

agagatctag cttcaggact cattggccct ctcctcatct gctacaaaga atctgtagat 1680

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cctggtacgt cgagcacggg gtcgccaggg gcgtcccctg ggacgtcctc aacaggctcg 2400

cccggtgcgt cacccggcac gtcgtccacg ggttcacctg gtagctcccc ttccgcgtcc 2460

actggcaccg ggcctggaac tccggggagc ggcacagcga gctcgtcgcc gggagcatcg 2520

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gccactggat caccagggag ctcgacacct tccggcgcaa cgggatcgcc cggagccagc 2700

ccgggtacgt caagcactgg ctcccctgcc tcgagctcag atcaagagga aatcgattat 2760

gatgatacca tatcagttga aatgaagaag gaagattttg acatttatga tgaggatgaa 2820

aatcagagcc cccgcagctt tcaaaagaaa acacgacact attttattgc tgcagtggag 2880

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cagcccttat accgtggaga actaaatgaa catttgggac tcctggggcc atatataaga 3060

gcagaagttg aagataatat catggtaact ttcagaaatc aggcctctcg tccctattcc 3120

ttctattcta gccttatttc ttatgaggaa gatcagaggc aaggagcaga acctagaaaa 3180

aactttgtca agcctaatga aaccaaaact tacttttgga aagtgcaaca tcatatggca 3240

cccactaaag atgagtttga ctgcaaagcc tgggcttatt tctctgatgt tgacctggaa 3300

aaagatgtgc actcaggcct gattggaccc cttctggtct gccacactaa cacactgaac 3360

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gagaccaaaa gctggtactt cactgaaaat atggaaagaa actgcagggc tccctgcaat 3480

atccagatgg aagatcccac ttttaaagag aattatcgct tccatgcaat caatggctac 3540

ataatggata cactacctgg cttagtaatg gctcaggatc aaaggattcg atggtatctg 3600

ctcagcatgg gcagcaatga aaacatccat tctattcatt tcagtggaca tgtgttcact 3660

gtacgaaaaa aagaggagta taaaatggca ctgtacaatc tctatccagg tgtttttgag 3720

acagtggaaa tgttaccatc caaagctgga atttggcggg tggaatgcct tattggcgag 3780

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aagacccagg gtgcccgtca gaagttctcc agcctctaca tctctcagtt tatcatcatg 4080

tatagtcttg atgggaagaa gtggcagact tatcgaggaa attccactgg aaccttaatg 4140

gtcttctttg gcaatgtgga ttcatctggg ataaaacaca atatttttaa ccctccaatt 4200

attgctcgat acatccgttt gcacccaact cattatagca ttcgcagcac tcttcgcatg 4260

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agtcaagatg gccatcagtg gactctcttt tttcagaatg gcaaagtaaa ggtttttcag 4620

ggaaatcaag actccttcac acctgtggtg aactctctag acccaccgtt actgactcgc 4680

taccttcgaa ttcaccccca gagttgggtg caccagattg ccctgaggat ggaggttctg 4740

ggctgcgagg cacaggacct ctacggcgcc ggatcacctg gggccgaaac ggccgaacaa 4800

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<210> 761
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 761

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

```
His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480
```

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
740 745 750

```
Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Gly Ala Pro Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
            820                 825                 830

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
        835                 840                 845

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
    850                 855                 860

Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
865                 870                 875                 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
                885                 890                 895

Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
            900                 905                 910

Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
        915                 920                 925

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
    930                 935                 940

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
945                 950                 955                 960

Ser Ala Pro Gly Ala Ser Ser Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020
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Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275
```

```
Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                1435                1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                1450                1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                1465                1470

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                1480                1485

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1490                1495                1500

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                1510                1515

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1520                1525                1530
```

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1580 1585 1590

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1595 1600 1605

Ser Pro Ala Thr Gly His His His His His His
1610 1615

<210> 762
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080
gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140
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tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560
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cccactaaag atgagtttga ctgcaaagcc tgggcttatt tctctgatgt tgacctggaa 3300

aaagatgtgc actcaggcct gattggaccc cttctggtct gccacactaa cacactgaac 3360

cctgctcatg ggagacaagt gacagtacag gaatttgctc tgtttttcac catctttgat 3420

gagaccaaaa gctggtactt cactgaaaat atggaaagaa actgcagggc tccctgcaat 3480

atccagatgg aagatcccac ttttaaagag aattatcgct tccatgcaat caatggctac 3540

ataatggata cactacctgg cttagtaatg gctcaggatc aaaggattcg atggtatctg 3600

ctcagcatgg gcagcaatga aaacatccat tctattcatt tcagtggaca tgtgttcact 3660

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ggctgcgagg cacaggacct ctacggcgcc ggatcacctg gggccgaaac ggccgaacaa 4800

aaactcatct cagaagagga tctgtcacct gcaaccggtc atcatcacca tcaccattga 4860

<210> 763
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 763
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

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Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525
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Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800
```

```
Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Gly
            820                 825                 830

Ala Pro Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
        835                 840                 845

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
    850                 855                 860

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
865                 870                 875                 880

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
                885                 890                 895

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
            900                 905                 910

Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
        915                 920                 925

Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro
    930                 935                 940

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr
945                 950                 955                 960

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
                965                 970                 975

Pro Gly Ala Ser Ser Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065
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Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320
```

```
Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                1435                1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                1450                1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                1465                1470

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                1480                1485

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1490                1495                1500

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                1510                1515

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1520                1525                1530

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1535                1540                1545

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1550                1555                1560

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1565                1570                1575
```

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1580 1585 1590

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1595 1600 1605

Ser Pro Ala Thr Gly His His His His His His
1610 1615

<210> 764
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

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gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

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ggctgcgagg cacaggacct ctacggcgcc ggatcacctg gggccgaaac ggccgaacaa 4800

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<211> 1619
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys

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Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
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305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln

580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
740 745 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
755 760 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
770 775 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785 790 795 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
805 810 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
820 825 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
835 840 845

Phe Thr Gln Pro Leu Tyr Arg Gly Ala Pro Thr Ser Glu Ser Ala Thr

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Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
865                 870                 875                 880

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
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Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
            900                 905                 910

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
        915                 920                 925

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro
    930                 935                 940

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro
945                 950                 955                 960

Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
                965                 970                 975

Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
            980                 985                 990

Thr Glu Pro Ser Glu Gly Ser Ala  Pro Gly Ala Ser Ser  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
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Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
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1115 1120 1125

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1130 1135 1140

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1145 1150 1155

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1160 1165 1170

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1175 1180 1185

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1190 1195 1200

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1205 1210 1215

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1220 1225 1230

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1235 1240 1245

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1250 1255 1260

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1265 1270 1275

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1280 1285 1290

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1295 1300 1305

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1310 1315 1320

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1325 1330 1335

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1340 1345 1350

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1355 1360 1365

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe

1370 1375 1380

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1385 1390 1395

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1400 1405 1410

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1415 1420 1425

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1430 1435 1440

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1445 1450 1455

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1460 1465 1470

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1475 1480 1485

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1490 1495 1500

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1505 1510 1515

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1520 1525 1530

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1580 1585 1590

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
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Ser Pro Ala Thr Gly His His His His His His
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<210> 766

<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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ggaaatcaag actccttcac acctgtggtg aactctctag acccaccgtt actgactcgc 4680

taccttcgaa ttcaccccca gagttgggtg caccagattg ccctgaggat ggaggttctg 4740

ggctgcgagg cacaggacct ctacggcgcc ggatcacctg gggccgaaac ggccgaacaa 4800

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<210> 767
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 767
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

```
Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365
```

```
Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640
```

```
Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910
```

```
Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gly Ala Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    1010                1015                1020

Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1025                1030                1035

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1040                1045                1050

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala
    1055                1060                1065

Ser Thr  Gly Thr Gly Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    1070                1075                1080

Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1085                1090                1095

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1100                1105                1110

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser
    1115                1120                1125

Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    1130                1135                1140

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1145                1150                1155

Ala Ser  Ser Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170
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```
Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425
```

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1430 1435 1440

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1445 1450 1455

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1460 1465 1470

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1475 1480 1485

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1490 1495 1500

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1505 1510 1515

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1520 1525 1530

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1580 1585 1590

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1595 1600 1605

Ser Pro Ala Thr Gly His His His His His His
1610 1615

<210> 768
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

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gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

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ggcgcaacgg gatcgcccgg agccagcccg ggtacgtcaa gcactggctc ccctgcctcg 3480

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ctcagcatgg gcagcaatga aaacatccat tctattcatt tcagtggaca tgtgttcact 3660

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taccttcgaa ttcaccccca gagttgggtg caccagattg ccctgaggat ggaggttctg 4740

ggctgcgagg cacaggacct ctacggcgcc ggatcacctg gggccgaaac ggccgaacaa 4800

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<210> 769
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 769

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

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Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415
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Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685
```

```
Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960
```

```
Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Gly Ala Pro Gly  Thr Pro Gly
    1010                1015                1020

Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1025                1030                1035

Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1040                1045                1050

Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly
    1055                1060                1065

Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly Pro Gly  Thr Pro Gly
    1070                1075                1080

Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1085                1090                1095

Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1100                1105                1110

Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly
    1115                1120                1125

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr
    1130                1135                1140

Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1145                1150                1155

Ser Thr  Gly Ser Pro Ala Ser  Ser Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215
```

```
Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                1435                1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                1450                1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                1465                1470
```

```
Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                 1480                 1485

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1490                 1495                 1500

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                 1510                 1515

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1520                 1525                 1530

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1535                 1540                 1545

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1550                 1555                 1560

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1565                 1570                 1575

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1580                 1585                 1590

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
    1595                 1600                 1605

Ser Pro  Ala Thr Gly His His  His His His His
    1610                 1615
```

<210> 770
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 770

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac      180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt      240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct      300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa      360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc      420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct      480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga      540
```

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<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 771
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

```
His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460
```

```
Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735
```

```
Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
        995                 1000                1005
```

```
Pro Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
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Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
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Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1040                1045                1050

Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1055                1060                1065

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1070                1075                1080

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Pro  Ala Gly Ser
    1085                1090                1095

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1100                1105                1110

Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1115                1120                1125

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1130                1135                1140

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ala Ser  Ser Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260
```

```
Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
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Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                1435                1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                1450                1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                1465                1470

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                1480                1485

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1490                1495                1500

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                1510                1515
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Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1520 1525 1530

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1580 1585 1590

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1595 1600 1605

Ser Pro Ala Thr Gly His His His His His His
1610 1615

<210> 772
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

agagatctag cttcaggact cattggccct ctcctcatct gctacaaaga atctgtagat 1680

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ataatggata cactacctgg cttagtaatg gctcaggatc aaaggattcg atggtatctg 3600

ctcagcatgg gcagcaatga aaacatccat tctattcatt tcagtggaca tgtgttcact 3660

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tatagtcttg atgggaagaa gtggcagact tatcgaggaa attccactgg aaccttaatg 4140

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<210> 773
<211> 1619
<212> PRT

<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 773
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His

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                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
```

515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
740 745 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
755 760 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
770 775 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys

785 790 795 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
805 810 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
820 825 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
835 840 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
850 855 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865 870 875 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
885 890 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
900 905 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
915 920 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
930 935 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945 950 955 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
965 970 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
980 985 990

Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Gly Ala
995 1000 1005

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala
1010 1015 1020

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
1025 1030 1035

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
1040 1045 1050

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly

1055 1060 1065

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala
1070 1075 1080

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
1085 1090 1095

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
1100 1105 1110

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
1115 1120 1125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala
1130 1135 1140

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Ala Pro
1145 1150 1155

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1160 1165 1170

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1175 1180 1185

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1190 1195 1200

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1205 1210 1215

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1220 1225 1230

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1235 1240 1245

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1250 1255 1260

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1265 1270 1275

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1280 1285 1290

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1295 1300 1305

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro

1310 1315 1320

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1325 1330 1335

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1340 1345 1350

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1355 1360 1365

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1370 1375 1380

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1385 1390 1395

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1400 1405 1410

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1415 1420 1425

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1430 1435 1440

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1445 1450 1455

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1460 1465 1470

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1475 1480 1485

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1490 1495 1500

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1505 1510 1515

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1520 1525 1530

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu

1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1580 1585 1590

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1595 1600 1605

Ser Pro Ala Thr Gly His His His His His His
1610 1615

<210> 774
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

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ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

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atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

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gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

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<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
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polypeptide

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20 25 30

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Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
        35                  40                  45

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
    50                  55                  60

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
65                  70                  75                  80

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
                85                  90                  95

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
            100                 105                 110

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
        115                 120                 125

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
    130                 135                 140

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
145                 150                 155                 160

Ser Thr Glu Glu Gly Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300
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Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575
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```
Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845
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Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110
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Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365
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Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                1435                1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                1450                1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                1465                1470

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                1480                1485

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1490                1495                1500

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                1510                1515

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1520                1525                1530

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1535                1540                1545

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1550                1555                1560

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1565                1570                1575

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1580                1585                1590

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
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Ser Pro  Ala Thr Gly His His  His His His His His
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<211> 4860
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<220>
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polynucleotide

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gccactagcg gctccgagac tccgggaact tccgagagcg ctacaccaga aagcggaccc 240

ggaaccagta ccgaacctag cgagggctct gctccgggca gcccagccgg ctctcctaca 300

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<211> 1619
<212> PRT
<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic polypeptide

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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
35 40 45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
50 55 60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65 70 75 80

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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
                85                  90                  95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
            100                 105                 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
        115                 120                 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
    130                 135                 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145                 150                 155                 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350
```

```
Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620
```

```
Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895
```

```
Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155
```

```
Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410
```

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1415 1420 1425

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1430 1435 1440

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1445 1450 1455

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1460 1465 1470

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1475 1480 1485

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1490 1495 1500

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1505 1510 1515

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1520 1525 1530

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1580 1585 1590

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
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Ser Pro Ala Thr Gly His His His His His His
1610 1615

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<211> 4860
<212> DNA
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<220>
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cccggaacgt catcgaccgg atcgccaggg agctcgacgc catcaggcgc aacaggatca 240
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<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 779
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Met Gln Ser Asp Leu Gly Gly Ala Pro Thr Ser Glu Ser Ala Thr Pro
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Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
35 40 45

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
50 55 60

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
65 70 75 80

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
85 90 95

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
100 105 110

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
115 120 125

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Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
    130                 135                 140

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala
145                 150                 155                 160

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ala Ser Ser Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400
```

```
Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670
```

```
Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940
```

```
Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200
```

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Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                 1210                 1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                 1225                 1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                 1240                 1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                 1255                 1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                 1270                 1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                 1285                 1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                 1300                 1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                 1315                 1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                 1330                 1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                 1345                 1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                 1360                 1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                 1375                 1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                 1390                 1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                 1405                 1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                 1420                 1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                 1435                 1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                 1450                 1455
```

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1460 1465 1470

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1475 1480 1485

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1490 1495 1500

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1505 1510 1515

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1520 1525 1530

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1580 1585 1590

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1595 1600 1605

Ser Pro Ala Thr Gly His His His His His His
1610 1615

<210> 780
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 780
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gccacatctg ggtcggaaac gccaggcaca agtgagtctg caactcccga gtccggacct 180

ggctccgagc ctgccactag cggctccgag actccgggaa cttccgagag cgctacacca 240

gaaagcggac ccggaaccag taccgaacct agcgagggct ctgctccggg cagcccagcc 300

ggctctccta catccacgga ggagggcact tccgaatccg ccaccccgga gtcagggcca 360

ggatctgaac ccgctacctc aggcagtgag acgccaggaa cgagcgagtc cgctacaccg 420

gagagtgggc cagggagccc tgctggatct cctacgtcca ctgaggaagg gtcaccagcg 480

ggctcgccca ccagcactga agaaggtgcc tcgagcgagc tgcctgtgga cgcaagattt 540

cctcctagag tgccaaaatc ttttccattc aacacctcag tcgtgtacaa aaagactctg 600

tttgtagaat tcacggatca ccttttcaac atcgctaagc caaggccacc ctggatgggt 660

ctgctaggtc ctaccatcca ggctgaggtt tatgatacag tggtcattac acttaagaac 720

atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag 780

ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct 840

ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

ggcctcattg gagccctact agtatgtaga gaagggagtc tggccaagga aaagacacag 1020

accttgcaca aatttatact acttttgct gtatttgatg aagggaaaag ttggcactca 1080

gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg 1200

aaatcagtct attggcatgt gattggaatg ggcaccactc ctgaagtgca ctcaatattc 1260

ctcgaaggtc acacatttct tgtgaggaac catcgccagg ctagcttgga aatctcgcca 1320

ataactttcc ttactgctca aacactcttg atggaccttg gacagtttct actgttttgt 1380

catatctctt cccaccaaca tgatggcatg gaagcttatg tcaaagtaga cagctgtcca 1440

gaggaacccc aactacgaat gaaaaataat gaagaagcgg aagactatga tgatgatctt 1500

actgattctg aaatggatgt ggtcaggttt gatgatgaca actctccttc ctttatccaa 1560

attcgctcag ttgccaagaa gcatcctaaa acttgggtac attacattgc tgctgaagag 1620

gaggactggg actatgctcc cttagtcctc gcccccgatg acagaagtta taaaagtcaa 1680

tatttgaaca atggccctca gcggattggt aggaagtaca aaaaagtccg atttatggca 1740

tacacagatg aaacctttaa gactcgtgaa gctattcagc atgaatcagg aatcttggga 1800

cctttacttt atggggaagt tggagacaca ctgttgatta tatttaagaa tcaagcaagc 1860

agaccatata acatctaccc tcacggaatc actgatgtcc gtcctttgta ttcaaggaga 1920

ttaccaaaag gtgtaaaaca tttgaaggat tttccaattc tgccaggaga aatattcaaa 1980

tataaatgga cagtgactgt agaagatggg ccaactaaat cagatcctcg gtgcctgacc 2040

cgctattact ctagtttcgt taatatggag agagatctag cttcaggact cattggccct 2100

ctcctcatct gctacaaaga atctgtagat caaagaggaa accagataat gtcagacaag 2160

aggaatgtca tcctgttttc tgtatttgat gagaaccgaa gctggtacct cacagagaat 2220

atacaacgct ttctccccaa tccagctgga gtgcagcttg aggatccaga gttccaagcc 2280

tccaacatca tgcacagcat caatggctat gtttttgata gtttgcagtt gtcagtttgt 2340

ttgcatgagg tggcatactg gtacattcta agcattggag cacagactga cttcctttct 2400

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gggtgccaca actcagactt tcggaacaga ggcatgaccg ccttactgaa ggtttctagt 2580

tgtgacaaga acactggtga ttattacgag gacagttatg aagatatttc agcatacttg 2640

ctgagtaaaa acaatgccat tgaaccaaga agcttctctc aaaacccacc agtcttgaaa 2700

cgccatcaac gggaaataac tcgtactact cttcagtcag atcaagagga aatcgattat 2760

gatgatacca tatcagttga aatgaagaag gaagattttg acatttatga tgaggatgaa 2820

aatcagagcc cccgcagctt tcaaaagaaa acacgacact attttattgc tgcagtggag 2880

aggctctggg attatgggat gagtagctcc ccacatgttc taagaaacag ggctcagagt 2940

ggcagtgtcc ctcagttcaa gaaagttgtt ttccaggaat ttactgatgg ctcctttact 3000

cagcccttat accgtggaga actaaatgaa catttgggac tcctggggcc atatataaga 3060

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aactttgtca agcctaatga aaccaaaact tacttttgga aagtgcaaca tcatatggca 3240

cccactaaag atgagtttga ctgcaaagcc tgggcttatt tctctgatgt tgacctggaa 3300

aaagatgtgc actcaggcct gattggaccc cttctggtct gccacactaa cacactgaac 3360

cctgctcatg ggagacaagt gacagtacag gaatttgctc tgtttttcac catctttgat 3420

gagaccaaaa gctggtactt cactgaaaat atggaaagaa actgcagggc tccctgcaat 3480

atccagatgg aagatcccac ttttaaagag aattatcgct tccatgcaat caatggctac 3540

ataatggata cactacctgg cttagtaatg gctcaggatc aaaggattcg atggtatctg 3600

ctcagcatgg gcagcaatga aaacatccat tctattcatt tcagtggaca tgtgttcact 3660

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aagacccagg gtgcccgtca gaagttctcc agcctctaca tctctcagtt tatcatcatg 4080

tatagtcttg atgggaagaa gtggcagact tatcgaggaa attccactgg aaccttaatg 4140

gtcttctttg gcaatgtgga ttcatctggg ataaaacaca atatttttaa ccctccaatt 4200

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ggaaatcaag actccttcac acctgtggtg aactctctag acccaccgtt actgactcgc 4680

taccttcgaa ttcaccccca gagttgggtg caccagattg ccctgaggat ggaggttctg 4740

ggctgcgagg cacaggacct ctacggcgcc ggatcacctg gggccgaaac ggccgaacaa 4800

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<210> 781
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 781

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Gly Ala Pro Gly Ser Ser Pro Ser Ala Ser
20 25 30

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
35 40 45

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
50 55 60

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly
65 70 75 80

Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
85 90 95

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
100 105 110

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr
115 120 125

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
130 135 140

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser
145 150 155 160

Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Glu Leu Pro Val
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr

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            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
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    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
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725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
835 840 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
885 890 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
900 905 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
915 920 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
930 935 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945 950 955 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
965 970 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
980 985 990

Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu

995 1000 1005

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1010 1015 1020

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1025 1030 1035

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1040 1045 1050

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1055 1060 1065

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1070 1075 1080

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1085 1090 1095

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1100 1105 1110

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1115 1120 1125

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1130 1135 1140

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1145 1150 1155

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1160 1165 1170

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1175 1180 1185

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1190 1195 1200

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1205 1210 1215

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1220 1225 1230

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1235 1240 1245

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His

1250 1255 1260

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1265 1270 1275

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1280 1285 1290

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1295 1300 1305

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1310 1315 1320

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1325 1330 1335

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1340 1345 1350

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1355 1360 1365

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1370 1375 1380

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1385 1390 1395

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1400 1405 1410

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1415 1420 1425

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1430 1435 1440

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1445 1450 1455

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1460 1465 1470

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1475 1480 1485

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1490 1495 1500

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile

1505 1510 1515

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1520 1525 1530

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1580 1585 1590

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1595 1600 1605

Ser Pro Ala Thr Gly His His His His His His
1610 1615

<210> 782
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 782
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ccctcagcat caacggggac gggaccgggg gcgtcaccag gaacgtcctc caccggctcg 180

ccgggtgcat cacccggaac gtcatcgacc ggatcgccag ggagctcgac gccatcaggc 240

gcaacaggat cacctggctc aagccctagc gcgtcaaccg gcacgggtcc gggtgcctcc 300

cctggcacgt ccagcaccgg atcacccgga tcgagcccat ccgcctcaac cggaaccgga 360

cccggtacac cagggtcggg aacagcctcc tcgtcaccag gctcctcaac cccctcggga 420

gccacgggtt cgcccggttc gtcaacgcct tccggagcaa ctggtagccc cggagcatcg 480

ccaggaactt cgagcacggg gtcgcccgcc tcgagcgagc tgcctgtgga cgcaagattt 540

cctcctagag tgccaaaatc ttttccattc aacacctcag tcgtgtacaa aaagactctg 600

tttgtagaat tcacggatca ccttttcaac atcgctaagc caaggccacc ctggatgggt 660

ctgctaggtc ctaccatcca ggctgaggtt tatgatacag tggtcattac acttaagaac 720

atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag 780

ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct 840

ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

ggcctcattg gagccctact agtatgtaga gaagggagtc tggccaagga aaagacacag 1020

accttgcaca aatttatact acttttgct gtatttgatg aagggaaaag ttggcactca 1080

gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg 1200

aaatcagtct attggcatgt gattggaatg ggcaccactc ctgaagtgca ctcaatattc 1260

ctcgaaggtc acacatttct tgtgaggaac catcgccagg ctagcttgga aatctcgcca 1320

ataactttcc ttactgctca aacactcttg atggaccttg gacagtttct actgttttgt 1380

catatctctt cccaccaaca tgatggcatg gaagcttatg tcaaagtaga cagctgtcca 1440

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<400> 783

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Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
        35                  40                  45

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
    50                  55                  60

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
65                  70                  75                  80

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
                85                  90                  95

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
            100                 105                 110

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
        115                 120                 125

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
    130                 135                 140

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
145                 150                 155                 160

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ala Ser Ser
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240
```

```
Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510
```

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Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780
```

```
Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
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Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
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Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
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Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
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Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
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Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
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Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
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Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
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Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
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Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
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Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
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Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
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Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
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Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
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Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
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Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
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Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
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Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
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Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
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Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
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Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
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Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
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Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                1510                1515

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1520                1525                1530

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1535                1540                1545

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1550                1555                1560
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Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
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Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
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actcagggag taaaatctct gcttaccagc atgtatgtga aggagttcct catctccagc 4560

agtcaagatg gccatcagtg gactctcttt tttcagaatg gcaaagtaaa ggtttttcag 4620

ggaaatcaag actccttcac acctgtggtg aactctctag acccaccgtt actgactcgc 4680

taccttcgaa ttcaccccca gagttgggtg caccagattg ccctgaggat ggaggttctg 4740

ggctgcgagg cacaggacct ctacggcgcc ggatcacctg gggccgaaac ggccgaacaa 4800

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<210> 785
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 785
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Gly Ser Ser
            20                  25                  30

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
        35                  40                  45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
    50                  55                  60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65                  70                  75                  80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
                85                  90                  95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
            100                 105                 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
        115                 120                 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
    130                 135                 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145                 150                 155                 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285
```

```
Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560
```

```
Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830
```

```
Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095
```

```
Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350
```

```
Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                1435                1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                1450                1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                1465                1470

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                1480                1485

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1490                1495                1500

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                1510                1515

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1520                1525                1530

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1535                1540                1545

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1550                1555                1560

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1565                1570                1575

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1580                1585                1590

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
    1595                1600                1605
```

Ser Pro Ala Thr Gly His His His His His His
1610 1615

<210> 786
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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ccaggttcgt caccctcagc atcaacgggg acgggaccgg gggcgtcacc aggaacgtcc 180

tccaccggct cgccgggtgc atcacccgga acgtcatcga ccggatcgcc agggagctcg 240

acgccatcag gcgcaacagg atcacctggc tcaagcccta gcgcgtcaac cggcacgggt 300

ccgggtgcct cccctggcac gtccagcacc ggatcacccg gatcgagccc atccgcctca 360

accggaaccg gacccggtac accagggtcg ggaacagcct cctcgtcacc aggctcctca 420

acccccctcgg gagccacggg ttcgcccggt tcgtcaacgc cttccggagc aactggtagc 480

cccggagcat cgccaggaac ttcgagcacg gggtcgcccg cctcgagcga cgcaagattt 540

cctcctagag tgccaaaatc ttttccattc aacacctcag tcgtgtacaa aaagactctg 600

tttgtagaat tcacggatca ccttttcaac atcgctaagc caaggccacc ctggatgggt 660

ctgctaggtc ctaccatcca ggctgaggtt tatgatacag tggtcattac acttaagaac 720

atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag 780

ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct 840

ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

ggcctcattg gagccctact agtatgtaga gaagggagtc tggccaagga aaagacacag 1020

accttgcaca aatttatact acttttgct gtatttgatg aagggaaaag ttggcactca 1080

gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg 1200

aaatcagtct attggcatgt gattggaatg ggcaccactc ctgaagtgca ctcaatattc 1260

ctcgaaggtc acacatttct tgtgaggaac catcgccagg ctagcttgga aatctcgcca 1320

ataactttcc ttactgctca aacactcttg atggaccttg gacagtttct actgttttgt 1380

catatctctt cccaccaaca tgatggcatg gaagcttatg tcaaagtaga cagctgtcca 1440

gaggaacccc aactacgaat gaaaaataat gaagaagcgg aagactatga tgatgatctt 1500

actgattctg aaatggatgt ggtcaggttt gatgatgaca actctccttc ctttatccaa 1560

attcgctcag ttgccaagaa gcatcctaaa acttgggtac attacattgc tgctgaagag 1620

gaggactggg actatgctcc cttagtcctc gcccccgatg acagaagtta taaaagtcaa 1680

tatttgaaca atggccctca gcggattggt aggaagtaca aaaaagtccg atttatggca 1740

tacacagatg aaacctttaa gactcgtgaa gctattcagc atgaatcagg aatcttggga 1800

cctttacttt atggggaagt tggagacaca ctgttgatta tatttaagaa tcaagcaagc 1860

agaccatata acatctaccc tcacggaatc actgatgtcc gtcctttgta ttcaaggaga 1920

ttaccaaaag gtgtaaaaca tttgaaggat tttccaattc tgccaggaga aatattcaaa 1980

tataaatgga cagtgactgt agaagatggg ccaactaaat cagatcctcg gtgcctgacc 2040

cgctattact ctagtttcgt taatatggag agagatctag cttcaggact cattggccct 2100

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gggtgccaca actcagactt tcggaacaga ggcatgaccg ccttactgaa ggtttctagt 2580

tgtgacaaga acactggtga ttattacgag gacagttatg aagatatttc agcatacttg 2640

ctgagtaaaa acaatgccat tgaaccaaga agcttctctc aaaacccacc agtcttgaaa 2700

cgccatcaac gggaaataac tcgtactact cttcagtcag atcaagagga aatcgattat 2760

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aatcagagcc cccgcagctt tcaaaagaaa acacgacact attttattgc tgcagtggag 2880

aggctctggg attatgggat gagtagctcc ccacatgttc taagaaacag ggctcagagt 2940

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cagcccttat accgtggaga actaaatgaa catttgggac tcctggggcc atatataaga 3060

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cccactaaag atgagtttga ctgcaaagcc tgggcttatt tctctgatgt tgacctggaa 3300

aaagatgtgc actcaggcct gattggaccc cttctggtct gccacactaa cacactgaac 3360

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gagaccaaaa gctggtactt cactgaaaat atggaaagaa actgcagggc tccctgcaat 3480

atccagatgg aagatcccac ttttaaagag aattatcgct tccatgcaat caatggctac 3540

ataatggata cactacctgg cttagtaatg gctcaggatc aaaggattcg atggtatctg 3600

ctcagcatgg gcagcaatga aaacatccat tctattcatt tcagtggaca tgtgttcact 3660

gtacgaaaaa aagaggagta taaaatggca ctgtacaatc tctatccagg tgtttttgag 3720

acagtggaaa tgttaccatc caaagctgga atttggcggg tggaatgcct tattggcgag 3780

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gtcttctttg gcaatgtgga ttcatctggg ataaaacaca atatttttaa ccctccaatt 4200

attgctcgat acatccgttt gcacccaact cattatagca ttcgcagcac tcttcgcatg 4260

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taccttcgaa ttcaccccca gagttgggtg caccagattg ccctgaggat ggaggttctg 4740

ggctgcgagg cacaggacct ctacggcgcc ggatcacctg gggccgaaac ggccgaacaa 4800

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<210> 787
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 787
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Gly Ala Pro Thr Ser Glu Ser Ala
35 40 45

Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu
50 55 60

```
Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
65                  70                  75                  80

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala
                85                  90                  95

Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
            100                 105                 110

Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr
        115                 120                 125

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr
    130                 135                 140

Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
145                 150                 155                 160

Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser
                165                 170                 175

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ala Ser Ser Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335
```

```
Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605
```

```
Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880
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Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140
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Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395
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Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1400 1405 1410

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1415 1420 1425

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1430 1435 1440

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1445 1450 1455

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1460 1465 1470

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1475 1480 1485

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1490 1495 1500

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1505 1510 1515

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1520 1525 1530

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1580 1585 1590

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1595 1600 1605

Ser Pro Ala Thr Gly His His His His His His
1610 1615

<210> 788
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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ggcgcgccaa catcagagag cgccaccccct gaaagtggtc ccgggagcga gccagccaca 180

tctgggtcgg aaacgccagg cacaagtgag tctgcaactc ccgagtccgg acctggctcc 240

gagcctgcca ctagcggctc cgagactccg ggaacttccg agagcgctac accagaaagc 300

ggacccggaa ccagtaccga acctagcgag ggctctgctc cgggcagccc agccggctct 360

cctacatcca cggaggaggg cacttccgaa tccgccaccc cggagtcagg gccaggatct 420

gaacccgcta cctcaggcag tgagacgcca ggaacgagcg agtccgctac accggagagt 480

gggccaggga gccctgctgg atctcctacg tccactgagg aagggtcacc agcgggctcg 540

cccaccagca ctgaagaagg tgcctcgagc aacacctcag tcgtgtacaa aaagactctg 600

tttgtagaat tcacggatca ccttttcaac atcgctaagc caaggccacc ctggatgggt 660

ctgctaggtc ctaccatcca ggctgaggtt tatgatacag tggtcattac acttaagaac 720

atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag 780

ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct 840

ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

ggcctcattg gagccctact agtatgtaga gaagggagtc tggccaagga aaagacacag 1020

accttgcaca aatttatact acttttgct gtatttgatg aagggaaaag ttggcactca 1080

gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg 1200

aaatcagtct attggcatgt gattggaatg ggcaccactc ctgaagtgca ctcaatattc 1260

ctcgaaggtc acacatttct tgtgaggaac catcgccagg ctagcttgga aatctcgcca 1320

ataactttcc ttactgctca aacactcttg atggaccttg gacagtttct actgttttgt 1380

catatctctt cccaccaaca tgatggcatg gaagcttatg tcaaagtaga cagctgtcca 1440

gaggaacccc aactacgaat gaaaaataat gaagaagcgg aagactatga tgatgatctt 1500

actgattctg aaatggatgt ggtcaggttt gatgatgaca actctccttc ctttatccaa 1560

attcgctcag ttgccaagaa gcatcctaaa acttgggtac attacattgc tgctgaagag 1620

gaggactggg actatgctcc cttagtcctc gcccccgatg acagaagtta taaaagtcaa 1680

tatttgaaca atggccctca gcggattggt aggaagtaca aaaaagtccg atttatggca 1740

tacacagatg aaacctttaa gactcgtgaa gctattcagc atgaatcagg aatcttggga 1800

cctttacttt atggggaagt tggagacaca ctgttgatta tatttaagaa tcaagcaagc 1860

agaccatata acatctaccc tcacggaatc actgatgtcc gtcctttgta ttcaaggaga 1920

ttaccaaaag gtgtaaaaca tttgaaggat tttccaattc tgccaggaga aatattcaaa 1980

tataaatgga cagtgactgt agaagatggg ccaactaaat cagatcctcg gtgcctgacc 2040

cgctattact ctagtttcgt taatatggag agagatctag cttcaggact cattggccct 2100

ctcctcatct gctacaaaga atctgtagat caaagaggaa accagataat gtcagacaag 2160

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ttcccattct caggagaaac tgtcttcatg tcgatggaaa acccaggtct atggattctg 2520

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tgtgacaaga acactggtga ttattacgag gacagttatg aagatatttc agcatacttg 2640

ctgagtaaaa acaatgccat tgaaccaaga agcttctctc aaaacccacc agtcttgaaa 2700

cgccatcaac gggaaataac tcgtactact cttcagtcag atcaagagga aatcgattat 2760

gatgatacca tatcagttga aatgaagaag gaagattttg acatttatga tgaggatgaa 2820

aatcagagcc cccgcagctt tcaaaagaaa acacgacact attttattgc tgcagtggag 2880

aggctctggg attatgggat gagtagctcc ccacatgttc taagaaacag ggctcagagt 2940

ggcagtgtcc ctcagttcaa gaaagttgtt ttccaggaat ttactgatgg ctcctttact 3000

cagcccttat accgtggaga actaaatgaa catttgggac tcctggggcc atatataaga 3060

gcagaagttg aagataatat catggtaact ttcagaaatc aggcctctcg tccctattcc 3120

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aactttgtca agcctaatga aaccaaaact tacttttgga aagtgcaaca tcatatggca 3240

cccactaaag atgagtttga ctgcaaagcc tgggcttatt tctctgatgt tgacctggaa 3300

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cctgctcatg ggagacaagt gacagtacag gaatttgctc tgtttttcac catctttgat 3420

gagaccaaaa gctggtactt cactgaaaat atggaaagaa actgcagggc tccctgcaat 3480

atccagatgg aagatcccac ttttaaagag aattatcgct tccatgcaat caatggctac 3540

ataatggata cactacctgg cttagtaatg gctcaggatc aaaggattcg atggtatctg 3600

ctcagcatgg gcagcaatga aaacatccat tctattcatt tcagtggaca tgtgttcact 3660

gtacgaaaaa aagaggagta taaaatggca ctgtacaatc tctatccagg tgtttttgag 3720

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ggctgcgagg cacaggacct ctacggcgcc ggatcacctg gggccgaaac ggccgaacaa 4800

aaactcatct cagaagagga tctgtcacct gcaaccggtc atcatcacca tcaccattga 4860

<210> 789
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 789
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Gly Ala Pro Gly Ser Ser Pro Ser
35 40 45

Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
50 55 60

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65 70 75 80

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro
85 90 95

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
100 105 110

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly

115 120 125

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser
130 135 140

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145 150 155 160

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
165 170 175

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
260 265 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
275 280 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
290 295 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305 310 315 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
325 330 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
340 345 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
355 360 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
370 375 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg

385 390 395 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
405 410 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
420 425 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
435 440 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
450 455 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465 470 475 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
485 490 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
500 505 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
515 520 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
530 535 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545 550 555 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
565 570 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
580 585 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
595 600 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
610 615 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
645 650 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr

660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
835 840 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
885 890 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
900 905 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
915 920 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro

930 935 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945 950 955 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
965 970 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
980 985 990

Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
995 1000 1005

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1010 1015 1020

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1025 1030 1035

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1040 1045 1050

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1055 1060 1065

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1070 1075 1080

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1085 1090 1095

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1100 1105 1110

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1115 1120 1125

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1130 1135 1140

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1145 1150 1155

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1160 1165 1170

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1175 1180 1185

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met

1190 1195 1200

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1205 1210 1215

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1220 1225 1230

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1235 1240 1245

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1250 1255 1260

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1265 1270 1275

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1280 1285 1290

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1295 1300 1305

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1310 1315 1320

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1325 1330 1335

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1340 1345 1350

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1355 1360 1365

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1370 1375 1380

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1385 1390 1395

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1400 1405 1410

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1415 1420 1425

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1430 1435 1440

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr

1445 1450 1455

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1460 1465 1470

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1475 1480 1485

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1490 1495 1500

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1505 1510 1515

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1520 1525 1530

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1580 1585 1590

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1595 1600 1605

Ser Pro Ala Thr Gly His His His His His His
1610 1615

<210> 790
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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ggcgcgccag gctccagccc ctccgcgagc acgggaaccg gaccaggttc gtcaccctca 180

gcatcaacgg ggacgggacc gggggcgtca ccaggaacgt cctccaccgg ctcgccgggt 240

gcatcacccg gaacgtcatc gaccggatcg ccagggagct cgacgccatc aggcgcaaca 300

ggatcacctg gctcaagccc tagcgcgtca accggcacgg gtccgggtgc ctcccctggc 360

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

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Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
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His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Gly Ala Pro Thr Ser Thr Glu Pro
    210                 215                 220

Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr
225                 230                 235                 240

Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
                245                 250                 255

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala
            260                 265                 270

Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
        275                 280                 285

Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
    290                 295                 300

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro
305                 310                 315                 320

Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
                325                 330                 335

Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
            340                 345                 350

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
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Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
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His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
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Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720
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Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
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Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
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Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990
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Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
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Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
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Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
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Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
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Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
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Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
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Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
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Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
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Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
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Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
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Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
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Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
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Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
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Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
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Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
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Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
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Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
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Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
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Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
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Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
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Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
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Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
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Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
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Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
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Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
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Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
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Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
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Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
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atccagatgg aagatcccac ttttaaagag aattatcgct tccatgcaat caatggctac 3540

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ctcagcatgg gcagcaatga aaacatccat tctattcatt tcagtggaca tgtgttcact 3660

gtacgaaaaa aagaggagta taaaatggca ctgtacaatc tctatccagg tgtttttgag 3720

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ggctgcgagg cacaggacct ctacggcgcc ggatcacctg gggccgaaac ggccgaacaa 4800

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<210> 793
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Gly Ala Pro Gly Ala Ser Pro Gly
210 215 220

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
225 230 235 240

Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
245 250 255

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
260 265 270

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
275 280 285

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
290 295 300

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
305 310 315 320

Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
325 330 335

Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
340 345 350

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Met Gln
355 360 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
370 375 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385 390 395 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
405 410 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
420 425 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
435 440 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
450 455 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465 470 475 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
485 490 495

```
Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765
```

```
Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035
```

```
Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290
```

```
Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                1435                1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                1450                1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                1465                1470

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                1480                1485

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1490                1495                1500

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                1510                1515

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1520                1525                1530

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1535                1540                1545
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Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1580 1585 1590

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1595 1600 1605

Ser Pro Ala Thr Gly His His His His His His
1610 1615

<210> 794
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

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ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttggg cgcgccaggt 660

gcctcgccgg gaacatcatc aactggttca cccgggtcat ccccctcggc ctcaaccggg 720

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ctcagcatgg gcagcaatga aaacatccat tctattcatt tcagtggaca tgtgttcact 3660

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<210> 795
<211> 1622
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 795

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Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270
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Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Ser Val Ala Gly Ala Pro Gly Ala Ser
    370                 375                 380

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
385                 390                 395                 400

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
                405                 410                 415

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser
            420                 425                 430

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
        435                 440                 445

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
    450                 455                 460

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser
465                 470                 475                 480

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
                485                 490                 495

Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
            500                 505                 510

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
        515                 520                 525

Lys Lys His Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu
    530                 535                 540
```

```
Asp Trp Asp Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr
545                 550                 555                 560

Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr
                565                 570                 575

Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg
            580                 585                 590

Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly
        595                 600                 605

Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg
    610                 615                 620

Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr
625                 630                 635                 640

Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile
                645                 650                 655

Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp
            660                 665                 670

Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser
        675                 680                 685

Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu
    690                 695                 700

Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met
705                 710                 715                 720

Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg
                725                 730                 735

Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala
            740                 745                 750

Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His
        755                 760                 765

Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu
    770                 775                 780

His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp
785                 790                 795                 800

Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val
                805                 810                 815
```

```
Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe
            820                 825                 830

Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser
        835                 840                 845

Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys
    850                 855                 860

Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser
865                 870                 875                 880

Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser
                885                 890                 895

Gln Asn Pro Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr
            900                 905                 910

Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser
        915                 920                 925

Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn
    930                 935                 940

Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala
945                 950                 955                 960

Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val
                965                 970                 975

Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val
            980                 985                 990

Val Phe Gln Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
        995                 1000                1005

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1010                1015                1020

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1025                1030                1035

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1040                1045                1050

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1055                1060                1065

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1070                1075                1080
```

```
Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1085                1090                1095

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1100                1105                1110

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1115                1120                1125

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1130                1135                1140

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1145                1150                1155

Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1160                1165                1170

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1175                1180                1185

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1190                1195                1200

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1205                1210                1215

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1220                1225                1230

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1235                1240                1245

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1250                1255                1260

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1265                1270                1275

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1280                1285                1290

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1295                1300                1305

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1310                1315                1320

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1325                1330                1335
```

```
Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1340                1345                1350

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1355                1360                1365

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1370                1375                1380

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1385                1390                1395

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1400                1405                1410

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1415                1420                1425

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1430                1435                1440

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1445                1450                1455

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1460                1465                1470

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1475                1480                1485

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1490                1495                1500

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1505                1510                1515

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1520                1525                1530

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1535                1540                1545

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    1550                1555                1560

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    1565                1570                1575

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    1580                1585                1590
```

Gly Ser Pro Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu
1595 1600 1605

Glu Asp Leu Ser Pro Ala Thr Gly His His His His His His
1610 1615 1620

<210> 796
<211> 4869
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgcctctgt agcaggcgcg 1140

ccaggtgcct cgccgggaac atcatcaact ggttcacccg ggtcatcccc ctcggcctca 1200

accgggacgg gtcccggctc atcccccagc gccagcactg gaacaggtcc tggcactcct 1260

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cctggggcat caccggggac gagctccacg gggtcccctg gagcgtcacc ggggacctcc 1560

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gctgaagagg aggactggga ctatgctccc ttagtcctcg cccccgatga cagaagttat 1680

aaaagtcaat atttgaacaa tggccctcag cggattggta ggaagtacaa aaaagtccga 1740

tttatggcat acacagatga aacctttaag actcgtgaag ctattcagca tgaatcagga 1800

atcttgggac ctttacttta tggggaagtt ggagacacac tgttgattat atttaagaat 1860

caagcaagca gaccatataa catctaccct cacggaatca ctgatgtccg tcctttgtat 1920

tcaaggagat taccaaaagg tgtaaaacat ttgaaggatt ttccaattct gccaggagaa 1980

atattcaaat ataaatggac agtgactgta gaagatgggc caactaaatc agatcctcgg 2040

tgcctgaccc gctattactc tagtttcgtt aatatggaga gagatctagc ttcaggactc 2100

attggccctc tcctcatctg ctacaaagaa tctgtagatc aaagaggaaa ccagataatg 2160

tcagacaaga ggaatgtcat cctgttttct gtatttgatg agaaccgaag ctggtacctc 2220

acagagaata tacaacgctt tctccccaat ccagctggag tgcagcttga ggatccagag 2280

ttccaagcct ccaacatcat gcacagcatc aatggctatg tttttgatag tttgcagttg 2340

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ccctgcaata tccagatgga agatcccact tttaaagaga attatcgctt ccatgcaatc 3540

aatggctaca taatggatac actacctggc ttagtaatgg ctcaggatca aaggattcga 3600

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cacggcatca agacccaggg tgcccgtcag aagttctcca gcctctacat ctctcagttt 4080

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acctggtctc cttcaaaagc tcgacttcac ctccaaggga ggagtaatgc ctggagacct 4440

caggtgaata atccaaaaga gtggctgcaa gtggacttcc agaagacaat gaaagtcaca 4500

ggagtaacta ctcagggagt aaaatctctg cttaccagca tgtatgtgaa ggagttcctc 4560

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gtttttcagg gaaatcaaga ctccttcaca cctgtggtga actctctaga cccaccgtta 4680

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gaggttctgg gctgcgaggc acaggacctc tacggcgccg gatcacctgg ggccgaaacg 4800

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<210> 797
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 797
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1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

```
Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320
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```
Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gly Ala Pro Thr Ser Thr
        435                 440                 445

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
    450                 455                 460

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
465                 470                 475                 480

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
                485                 490                 495

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
            500                 505                 510

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        515                 520                 525

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr
    530                 535                 540

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
545                 550                 555                 560

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
                565                 570                 575

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser
            580                 585                 590
```

```
Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860
```

```
Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125
```

```
Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380
```

```
Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
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Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
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Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                1435                1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                1450                1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                1465                1470

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                1480                1485

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1490                1495                1500

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                1510                1515

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1520                1525                1530

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1535                1540                1545

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1550                1555                1560

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1565                1570                1575

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1580                1585                1590

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
    1595                1600                1605

Ser Pro  Ala Thr Gly His His  His His His His His
    1610                1615
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<210> 798
<211> 4860
<212> DNA

<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattggcg cgccaaccag tacggagccg tccgagggga gcgcaccagg aagcccggct 1380

gggagcccga cttctaccga agagggtaca tctaccgaac caagtgaagg ttcagcacca 1440

ggcacctcaa cagaaccctc tgagggctcg gcgcctggta caagtgagtc cgccacccca 1500

gaatccgggc ctgggacaag cacagaacct tcggaaggga gtgcccctgg aacatccgaa 1560

tcggcaaccc cagaatcagg gccaggatct gagcccgcga cttcgggctc cgagacgcct 1620

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tcggcgaccc cggagagcgg gccaggtgcc tcgagccagc atgaatcagg aatcttggga 1800

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ctcagcatgg gcagcaatga aaacatccat tctattcatt tcagtggaca tgtgttcact 3660

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ggctgcgagg cacaggacct ctacggcgcc ggatcacctg gggccgaaac ggccgaacaa 4800

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<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

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Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365
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Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gly Ala Pro Gly Ala Ser
        435                 440                 445

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
    450                 455                 460

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
465                 470                 475                 480

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser
                485                 490                 495

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
            500                 505                 510

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
        515                 520                 525

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser
    530                 535                 540

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
545                 550                 555                 560

Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
                565                 570                 575

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640
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Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910
```

```
Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170
```

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Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425
```

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1430 1435 1440

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1445 1450 1455

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1460 1465 1470

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1475 1480 1485

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1490 1495 1500

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1505 1510 1515

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1520 1525 1530

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1580 1585 1590

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1595 1600 1605

Ser Pro Ala Thr Gly His His His His His His
1610 1615

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<211> 4860
<212> DNA
<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

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gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

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ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

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gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

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atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

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agaccatata acatctaccc tcacggaatc actgatgtcc gtcctttgta ttcaaggaga 1920

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tataaatgga cagtgactgt agaagatggg ccaactaaat cagatcctcg gtgcctgacc 2040

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<210> 801
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 801

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser

145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr

420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala

690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
740 745 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
755 760 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
770 775 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785 790 795 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
805 810 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
820 825 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
835 840 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
850 855 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865 870 875 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
885 890 895

Ser Tyr Glu Glu Asp Gln Gly Ala Pro Thr Ser Glu Ser Ala Thr Pro
900 905 910

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
915 920 925

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
930 935 940

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
945 950 955 960

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro

965 970 975

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
980 985 990

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
995 1000 1005

Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
1010 1015 1020

Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1025 1030 1035

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Arg
1040 1045 1050

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1055 1060 1065

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1070 1075 1080

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1085 1090 1095

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1100 1105 1110

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1115 1120 1125

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1130 1135 1140

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1145 1150 1155

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1160 1165 1170

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1175 1180 1185

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1190 1195 1200

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1205 1210 1215

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn

1220 1225 1230

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1235 1240 1245

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1250 1255 1260

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1265 1270 1275

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1280 1285 1290

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1295 1300 1305

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1310 1315 1320

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1325 1330 1335

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1340 1345 1350

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1355 1360 1365

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1370 1375 1380

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1385 1390 1395

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1400 1405 1410

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1415 1420 1425

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1430 1435 1440

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1445 1450 1455

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1460 1465 1470

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val

1475 1480 1485

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1490 1495 1500

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1505 1510 1515

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1520 1525 1530

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1580 1585 1590

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1595 1600 1605

Ser Pro Ala Thr Gly His His His His His His
1610 1615

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<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

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<210> 803
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

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His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480
```

```
Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750
```

```
Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr
            900                 905                 910

Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
        915                 920                 925

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
    930                 935                 940

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
945                 950                 955                 960

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
                965                 970                 975

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
            980                 985                 990

Pro Gly Thr Ser Ser Thr Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
        995                 1000                1005

Ser Thr  Gly Ser Pro Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly
    1010                1015                1020
```

```
Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly
    1025                1030                1035

Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser Pro Ala  Ser Ser Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275
```

```
Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                1435                1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                1450                1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                1465                1470

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                1480                1485

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1490                1495                1500

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                1510                1515

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1520                1525                1530
```

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1580 1585 1590

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1595 1600 1605

Ser Pro Ala Thr Gly His His His His His His
1610 1615

<210> 804
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

agagatctag cttcaggact cattggccct ctcctcatct gctacaaaga atctgtagat 1680

caaagaggaa accagataat gtcagacaag aggaatgtca tcctgttttc tgtatttgat 1740

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cttcagtcag atcaagagga aatcgattat gatgatacca tatcagttga aatgaagaag 2340

gaagattttg acatttatga tgaggatgaa aatcagagcc cccgcagctt tcaaaagaaa 2400

acacgacact attttattgc tgcagtggag aggctctggg attatgggat gagtagctcc 2460

ccacatgttc taagaaacag ggctcagagt ggcagtgtcc ctcagttcaa gaaagttgtt 2520

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cccactaaag atgagtttga ctgcaaagcc tgggcttatt tctctgatgt tgacctggaa 3300

aaagatgtgc actcaggcct gattggaccc cttctggtct gccacactaa cacactgaac 3360

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gagaccaaaa gctggtactt cactgaaaat atggaaagaa actgcagggc tccctgcaat 3480

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ataatggata cactacctgg cttagtaatg gctcaggatc aaaggattcg atggtatctg 3600

ctcagcatgg gcagcaatga aaacatccat tctattcatt tcagtggaca tgtgttcact 3660

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gtcttctttg gcaatgtgga ttcatctggg ataaaacaca atatttttaa ccctccaatt 4200

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ggctgcgagg cacaggacct ctacggcgcc ggatcacctg gggccgaaac ggccgaacaa 4800

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<210> 805
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 805

```
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255
```

```
Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525
```

```
Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800
```

```
Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Gly Ala Pro Thr
            900                 905                 910

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr
        915                 920                 925

Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
    930                 935                 940

Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr
945                 950                 955                 960

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro
                965                 970                 975

Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
            980                 985                 990

Gly Pro Gly Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser
        995                 1000                1005

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    1010                1015                1020

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1025                1030                1035

Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    1040                1045                1050

Pro Gly  Ala Ser Ser Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065
```

```
Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320
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```
Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                1435                1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                1450                1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                1465                1470

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                1480                1485

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1490                1495                1500

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                1510                1515

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1520                1525                1530

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1535                1540                1545

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1550                1555                1560

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1565                1570                1575
```

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1580 1585 1590

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1595 1600 1605

Ser Pro Ala Thr Gly His His His His His His
1610 1615

<210> 806
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 806
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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

agagatctag cttcaggact cattggccct ctcctcatct gctacaaaga atctgtagat 1680

caaagaggaa accagataat gtcagacaag aggaatgtca tcctgttttc tgtatttgat 1740

gagaaccgaa gctggtacct cacagagaat atacaacgct ttctccccaa tccagctgga 1800

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gtttttgata gtttgcagtt gtcagtttgt ttgcatgagg tggcatactg gtacattcta 1920

agcattggag cacagactga cttccttttct gtcttcttct ctggatatac cttcaaacac 1980

aaaatggtct atgaagacac actcacccta ttcccattct caggagaaac tgtcttcatg 2040

tcgatggaaa acccaggtct atggattctg gggtgccaca actcagactt tcggaacaga 2100

ggcatgaccg ccttactgaa ggtttctagt tgtgacaaga acactggtga ttattacgag 2160

gacagttatg aagatatttc agcatacttg ctgagtaaaa acaatgccat tgaaccaaga 2220

agcttctctc aaaacccacc agtcttgaaa cgccatcaac gggaaataac tcgtactact 2280

cttcagtcag atcaagagga aatcgattat gatgatacca tatcagttga aatgaagaag 2340

gaagattttg acatttatga tgaggatgaa aatcagagcc cccgcagctt tcaaaagaaa 2400

acacgacact attttattgc tgcagtggag aggctctggg attatgggat gagtagctcc 2460

ccacatgttc taagaaacag ggctcagagt ggcagtgtcc ctcagttcaa gaaagttgtt 2520

ttccaggaat ttactgatgg ctcctttact cagcccttat accgtggaga actaaatgaa 2580

catttgggac tcctggggcc atatataaga gcagaagttg aagataatat catggtaact 2640

ttcagaaatc aggcctctcg tccctattcc ttctattcta gccttatttc ttatgaggaa 2700

gatcagaggc aaggagcaga acctggcgcg ccaacgtccg aaagtgctac ccctgagtca 2760

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aactttgtca agcctaatga aaccaaaact tacttttgga aagtgcaaca tcatatggca 3240

cccactaaag atgagtttga ctgcaaagcc tgggcttatt tctctgatgt tgacctggaa 3300

aaagatgtgc actcaggcct gattggaccc cttctggtct gccacactaa cacactgaac 3360

cctgctcatg ggagacaagt gacagtacag gaatttgctc tgtttttcac catctttgat 3420

gagaccaaaa gctggtactt cactgaaaat atggaaagaa actgcagggc tccctgcaat 3480

atccagatgg aagatcccac ttttaaagag aattatcgct tccatgcaat caatggctac 3540

ataatggata cactacctgg cttagtaatg gctcaggatc aaaggattcg atggtatctg 3600

ctcagcatgg gcagcaatga aaacatccat tctattcatt tcagtggaca tgtgttcact 3660

gtacgaaaaa aagaggagta taaaatggca ctgtacaatc tctatccagg tgtttttgag 3720

acagtggaaa tgttaccatc caaagctgga atttggcggg tggaatgcct tattggcgag 3780

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aagacccagg gtgcccgtca gaagttctcc agcctctaca tctctcagtt tatcatcatg 4080

tatagtcttg atgggaagaa gtggcagact tatcgaggaa attccactgg aaccttaatg 4140

gtcttctttg gcaatgtgga ttcatctggg ataaaacaca atatttttaa ccctccaatt 4200

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ggctgcgagg cacaggacct ctacggcgcc ggatcacctg gggccgaaac ggccgaacaa 4800

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<210> 807
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 807
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1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300
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Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575
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Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845
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Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Gly Ala Pro Gly
            900                 905                 910

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly
        915                 920                 925

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
    930                 935                 940

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
945                 950                 955                 960

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser
                965                 970                 975

Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
            980                 985                 990

Gly Ser Pro Gly Ala Ser Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly
        995                 1000                1005

Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser Pro Gly  Ser Ser Thr
    1010                1015                1020

Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser Ser Thr  Pro Ser Gly
    1025                1030                1035

Ala Thr  Gly Ser Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1040                1045                1050

Ser Pro  Ala Ser Ser Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110
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```
Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365
```

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Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                1435                1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                1450                1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                1465                1470

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                1480                1485

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1490                1495                1500

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                1510                1515

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1520                1525                1530

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1535                1540                1545

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1550                1555                1560

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1565                1570                1575

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1580                1585                1590

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
    1595                1600                1605

Ser Pro  Ala Thr Gly His His  His His His His His
    1610                1615
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<210> 808
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val

85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
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Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe

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Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
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Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
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Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
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Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
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Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
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Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
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Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
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Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
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Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
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Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
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625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
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Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Thr Ser Glu
740 745 750

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
755 760 765

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
770 775 780

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785 790 795 800

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900 905 910

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945 950 955 960

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Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
995 1000 1005

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Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Ala Ser
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1040 1045 1050

Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
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Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp
1070 1075 1080

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Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser
1100 1105 1110

Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro
1115 1120 1125

Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe
1130 1135 1140

Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
1145 1150 1155

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val

1160 1165 1170

Thr Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser
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Leu Ile Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg
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Lys Asn Phe Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys
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Val Gln His His Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys
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Ser Gly Leu Ile Gly Pro Leu Leu Val Cys His Thr Asn Thr Leu
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Arg Ile Arg Trp Tyr Leu Leu Ser Met Gly Ser Asn Glu Asn Ile
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His Ser Ile His Phe Ser Gly His Val Phe Thr Val Arg Lys Lys
1355 1360 1365

Glu Glu Tyr Lys Met Ala Leu Tyr Asn Leu Tyr Pro Gly Val Phe
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Glu Thr Val Glu Met Leu Pro Ser Lys Ala Gly Ile Trp Arg Val
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Glu Cys Leu Ile Gly Glu His Leu His Ala Gly Met Ser Thr Leu
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Ser Gly His Ile Arg Asp Phe Gln Ile Thr Ala Ser Gly Gln Tyr
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1805 1810 1815

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1820 1825 1830

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr
1835 1840 1845

Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
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Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
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<210> 811
<211> 1759
<212> PRT
<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic polypeptide

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20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

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Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
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Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
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Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525
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Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
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Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
            740                 745                 750

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
        755                 760                 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
    770                 775                 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785                 790                 795                 800
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Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
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Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            820                 825                 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
        835                 840                 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    850                 855                 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865                 870                 875                 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065
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Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
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Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320
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Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
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Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                1435                1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                1450                1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                1465                1470

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                1480                1485

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1490                1495                1500

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                1510                1515

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1520                1525                1530

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1535                1540                1545

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
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Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1565                1570                1575
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Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Thr Ser
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Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala
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Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1610                 1615                 1620

Ser Gly  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1625                 1630                 1635

Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu
    1640                 1645                 1650

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser
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Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
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Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
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Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser
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Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro
    1715                 1720                 1725

Ser Glu  Gly Ser Ala Pro Gly  Ala Ser Ser Glu Gln  Lys Leu Ile
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Ser Glu  Glu Asp Leu Ser Pro  Ala Thr Gly His His  His His His
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<211> 1759
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

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Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365
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Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640
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Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Thr Pro Gly
            740                 745                 750

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
        755                 760                 765

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    770                 775                 780

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
785                 790                 795                 800

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
                805                 810                 815

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            820                 825                 830

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
        835                 840                 845

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
    850                 855                 860

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
865                 870                 875                 880

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910
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Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170
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```
Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425
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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                 1435                 1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                 1450                 1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                 1465                 1470

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                 1480                 1485

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1490                 1495                 1500

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                 1510                 1515

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1520                 1525                 1530

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1535                 1540                 1545

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1550                 1555                 1560

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1565                 1570                 1575

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Thr Ser
    1580                 1585                 1590

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala
    1595                 1600                 1605

Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1610                 1615                 1620

Ser Gly  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1625                 1630                 1635

Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu
    1640                 1645                 1650

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser
    1655                 1660                 1665

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1670                 1675                 1680
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Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1685                 1690                 1695

Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser
    1700                 1705                 1710

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro
    1715                 1720                 1725

Ser Glu  Gly Ser Ala Pro Gly  Ala Ser Ser Glu Gln  Lys Leu Ile
    1730                 1735                 1740

Ser Glu  Glu Asp Leu Ser Pro  Ala Thr Gly His His  His His His
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His
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<211> 5280
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

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<210> 815
<211> 1759
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

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His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480
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Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Thr Pro Gly
            740                 745                 750
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Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
        755                 760                 765

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    770                 775                 780

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
785                 790                 795                 800

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
                805                 810                 815

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            820                 825                 830

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
        835                 840                 845

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
    850                 855                 860

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
865                 870                 875                 880

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020
```

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Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275
```

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Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                1435                1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                1450                1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                1465                1470

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                1480                1485

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1490                1495                1500

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                1510                1515

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1520                1525                1530
```

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Pro Thr Ser
1580 1585 1590

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala
1595 1600 1605

Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
1610 1615 1620

Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
1625 1630 1635

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu
1640 1645 1650

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser
1655 1660 1665

Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
1670 1675 1680

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
1685 1690 1695

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser
1700 1705 1710

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro
1715 1720 1725

Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Glu Gln Lys Leu Ile
1730 1735 1740

Ser Glu Glu Asp Leu Ser Pro Ala Thr Gly His His His His His
1745 1750 1755

His

<210> 816
<211> 5280

<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 816
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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

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caaagaggaa accagataat gtcagacaag aggaatgtca tcctgttttc tgtatttgat 1740

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aggctctggg attatgggat gagtagctcc ccacatgttc taagaaacag ggctcagagt 2940

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gagccagcaa ccagcgggtc cgagactccc gggacctctg agtctgccac cccagagagc 5160

ggacccggca cttcaaccga gccctccgaa ggatcagcac caggtgcctc gagcgaacaa 5220

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<210> 817
<211> 1759
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 817
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys

```
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
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305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
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580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
740 745 750

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
755 760 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
770 775 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785 790 795 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
805 810 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
820 825 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
835 840 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly

850 855 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865 870 875 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
885 890 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
900 905 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
915 920 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
930 935 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945 950 955 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
965 970 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
980 985 990

Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
995 1000 1005

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1010 1015 1020

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1025 1030 1035

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1040 1045 1050

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1055 1060 1065

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1070 1075 1080

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1085 1090 1095

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1100 1105 1110

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr

1115 1120 1125

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1130 1135 1140

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1145 1150 1155

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1160 1165 1170

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1175 1180 1185

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1190 1195 1200

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1205 1210 1215

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1220 1225 1230

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1235 1240 1245

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1250 1255 1260

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1265 1270 1275

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1280 1285 1290

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1295 1300 1305

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1310 1315 1320

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1325 1330 1335

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1340 1345 1350

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1355 1360 1365

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe

1370 1375 1380

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1385 1390 1395

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1400 1405 1410

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1415 1420 1425

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1430 1435 1440

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1445 1450 1455

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1460 1465 1470

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1475 1480 1485

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1490 1495 1500

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1505 1510 1515

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1520 1525 1530

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Pro Thr Ser
1580 1585 1590

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala
1595 1600 1605

Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
1610 1615 1620

Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro

1625 1630 1635

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu
1640 1645 1650

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser
1655 1660 1665

Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
1670 1675 1680

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
1685 1690 1695

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser
1700 1705 1710

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro
1715 1720 1725

Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Glu Gln Lys Leu Ile
1730 1735 1740

Ser Glu Glu Asp Leu Ser Pro Ala Thr Gly His His His His His
1745 1750 1755

His

<210> 818
<211> 5280
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660
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cgccatcaag ctgaaataac tcgtactact cttcagtcag atcaagagga aatcgattat 2760

gatgatacca tatcagttga aatgaagaag gaagattttg acatttatga tgaggatgaa 2820

aatcagagcc cccgcagctt tcaaaagaaa acacgacact attttattgc tgcagtggag 2880

aggctctggg attatgggat gagtagctcc ccacatgttc taagaaacag ggctcagagt 2940

ggcagtgtcc ctcagttcaa gaaagttgtt ttccaggaat ttactgatgg ctcctttact 3000

cagcccttat accgtggaga actaaatgaa catttgggac tcctggggcc atatataaga 3060

gcagaagttg aagataatat catggtaact ttcagaaatc aggcctctcg tccctattcc 3120

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aactttgtca agcctaatga aaccaaaact tacttttgga aagtgcaaca tcatatggca 3240

cccactaaag atgagtttga ctgcaaagcc tgggcttatt tctctgatgt tgacctggaa 3300

aaagatgtgc actcaggcct gattggaccc cttctggtct gccacactaa cacactgaac 3360

cctgctcatg ggagacaagt gacagtacag gaatttgctc tgtttttcac catctttgat 3420

gagaccaaaa gctggtactt cactgaaaat atggaaagaa actgcagggc tccctgcaat 3480

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ctcagcatgg gcagcaatga aaacatccat tctattcatt tcagtggaca tgtgttcact 3660

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gagccagcaa ccagcgggtc cgagactccc gggacctctg agtctgccac cccagagagc 5160

ggacccggca cttcaaccga gccctccgaa ggatcagcac caggtgcctc gagcgaacaa 5220

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<210> 819
<211> 1903
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 819

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

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Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415
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Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685
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Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Thr Ser Glu
            740                 745                 750

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
        755                 760                 765

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
    770                 775                 780

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
785                 790                 795                 800

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
                805                 810                 815

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
            820                 825                 830

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
        835                 840                 845

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
    850                 855                 860

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
865                 870                 875                 880

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
                885                 890                 895

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
            900                 905                 910

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        915                 920                 925

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
    930                 935                 940

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
945                 950                 955                 960
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Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
               965                 970                 975

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
           980                 985                 990

Glu Pro Ser Glu Gly Ser Ala Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
       995                1000                1005

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1010                1015                1020

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Ala Ser
    1025                1030                1035

Ser Pro  Pro Val Leu Lys Arg  His Gln Ala Glu Ile  Thr Arg Thr
    1040                1045                1050

Thr Leu  Gln Ser Asp Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile
    1055                1060                1065

Ser Val  Glu Met Lys Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp
    1070                1075                1080

Glu Asn  Gln Ser Pro Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr
    1085                1090                1095

Phe Ile  Ala Ala Val Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser
    1100                1105                1110

Ser Pro  His Val Leu Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro
    1115                1120                1125

Gln Phe  Lys Lys Val Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe
    1130                1135                1140

Thr Gln  Pro Leu Tyr Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu
    1145                1150                1155

Leu Gly  Pro Tyr Ile Arg Ala  Glu Val Glu Asp Asn  Ile Met Val
    1160                1165                1170

Thr Phe  Arg Asn Gln Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser
    1175                1180                1185

Leu Ile  Ser Tyr Glu Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg
    1190                1195                1200

Lys Asn  Phe Val Lys Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys
    1205                1210                1215
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Val Gln  His His Met Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys
    1220                 1225                 1230

Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu Glu Lys  Asp Val His
    1235                 1240                 1245

Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys His Thr  Asn Thr Leu
    1250                 1255                 1260

Asn Pro  Ala His Gly Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu
    1265                 1270                 1275

Phe Phe  Thr Ile Phe Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu
    1280                 1285                 1290

Asn Met  Glu Arg Asn Cys Arg  Ala Pro Cys Asn Ile  Gln Met Glu
    1295                 1300                 1305

Asp Pro  Thr Phe Lys Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly
    1310                 1315                 1320

Tyr Ile  Met Asp Thr Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln
    1325                 1330                 1335

Arg Ile  Arg Trp Tyr Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile
    1340                 1345                 1350

His Ser  Ile His Phe Ser Gly  His Val Phe Thr Val  Arg Lys Lys
    1355                 1360                 1365

Glu Glu  Tyr Lys Met Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe
    1370                 1375                 1380

Glu Thr  Val Glu Met Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val
    1385                 1390                 1395

Glu Cys  Leu Ile Gly Glu His  Leu His Ala Gly Met  Ser Thr Leu
    1400                 1405                 1410

Phe Leu  Val Tyr Ser Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala
    1415                 1420                 1425

Ser Gly  His Ile Arg Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr
    1430                 1435                 1440

Gly Gln  Trp Ala Pro Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser
    1445                 1450                 1455

Ile Asn  Ala Trp Ser Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val
    1460                 1465                 1470
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Asp Leu  Leu Ala Pro Met Ile  Ile His Gly Ile Lys  Thr Gln Gly
    1475                1480                1485

Ala Arg  Gln Lys Phe Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile
    1490                1495                1500

Met Tyr  Ser Leu Asp Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn
    1505                1510                1515

Ser Thr  Gly Thr Leu Met Val  Phe Phe Gly Asn Val  Asp Ser Ser
    1520                1525                1530

Gly Ile  Lys His Asn Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr
    1535                1540                1545

Ile Arg  Leu His Pro Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg
    1550                1555                1560

Met Glu  Leu Met Gly Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu
    1565                1570                1575

Gly Met  Glu Ser Lys Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser
    1580                1585                1590

Ser Tyr  Phe Thr Asn Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala
    1595                1600                1605

Arg Leu  His Leu Gln Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val
    1610                1615                1620

Asn Asn  Pro Lys Glu Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met
    1625                1630                1635

Lys Val  Thr Gly Val Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr
    1640                1645                1650

Ser Met  Tyr Val Lys Glu Phe  Leu Ile Ser Ser Ser  Gln Asp Gly
    1655                1660                1665

His Gln  Trp Thr Leu Phe Phe  Gln Asn Gly Lys Val  Lys Val Phe
    1670                1675                1680

Gln Gly  Asn Gln Asp Ser Phe  Thr Pro Val Val Asn  Ser Leu Asp
    1685                1690                1695

Pro Pro  Leu Leu Thr Arg Tyr  Leu Arg Ile His Pro  Gln Ser Trp
    1700                1705                1710

Val His  Gln Ile Ala Leu Arg  Met Glu Val Leu Gly  Cys Glu Ala
    1715                1720                1725
```

Gln Asp Leu Tyr Gly Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly
1730 1735 1740

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
1745 1750 1755

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
1760 1765 1770

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr
1775 1780 1785

Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
1790 1795 1800

Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
1805 1810 1815

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
1820 1825 1830

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr
1835 1840 1845

Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
1850 1855 1860

Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
1865 1870 1875

Gly Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
1880 1885 1890

Pro Ala Thr Gly His His His His His His
1895 1900

<210> 820
<211> 5712
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

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20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

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Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
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Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320
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Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590
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Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
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Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Pro Gly Ala Ser
            740                 745                 750

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
        755                 760                 765

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
    770                 775                 780

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro
785                 790                 795                 800

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
                805                 810                 815

Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
            820                 825                 830

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
        835                 840                 845

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
    850                 855                 860
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Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
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Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
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Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
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Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
        915                 920                 925

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
    930                 935                 940

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
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Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
                965                 970                 975

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
            980                 985                 990

Pro Gly Thr Ser Ser Thr Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser
        995                 1000                1005

Thr Gly  Thr Gly Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
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Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Ala Ser
    1025                1030                1035

Ser Pro  Pro Val Leu Lys Arg  His Gln Ala Glu Ile  Thr Arg Thr
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Thr Leu  Gln Ser Asp Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile
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Ser Val  Glu Met Lys Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp
    1070                1075                1080

Glu Asn  Gln Ser Pro Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr
    1085                1090                1095

Phe Ile  Ala Ala Val Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser
    1100                1105                1110

Ser Pro  His Val Leu Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro
    1115                1120                1125
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Gln Phe  Lys Lys Val Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe
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Thr Gln  Pro Leu Tyr Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu
    1145                1150                1155

Leu Gly  Pro Tyr Ile Arg Ala  Glu Val Glu Asp Asn  Ile Met Val
    1160                1165                1170

Thr Phe  Arg Asn Gln Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser
    1175                1180                1185

Leu Ile  Ser Tyr Glu Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg
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Lys Asn  Phe Val Lys Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys
    1205                1210                1215

Val Gln  His His Met Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys
    1220                1225                1230

Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu Glu Lys  Asp Val His
    1235                1240                1245

Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys His Thr  Asn Thr Leu
    1250                1255                1260

Asn Pro  Ala His Gly Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu
    1265                1270                1275

Phe Phe  Thr Ile Phe Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu
    1280                1285                1290

Asn Met  Glu Arg Asn Cys Arg  Ala Pro Cys Asn Ile  Gln Met Glu
    1295                1300                1305

Asp Pro  Thr Phe Lys Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly
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Tyr Ile  Met Asp Thr Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln
    1325                1330                1335

Arg Ile  Arg Trp Tyr Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile
    1340                1345                1350

His Ser  Ile His Phe Ser Gly  His Val Phe Thr Val  Arg Lys Lys
    1355                1360                1365

Glu Glu  Tyr Lys Met Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe
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Glu Thr  Val Glu Met Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val
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Glu Cys  Leu Ile Gly Glu His  Leu His Ala Gly Met  Ser Thr Leu
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Phe Leu  Val Tyr Ser Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala
    1415                1420                1425

Ser Gly  His Ile Arg Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr
    1430                1435                1440

Gly Gln  Trp Ala Pro Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser
    1445                1450                1455

Ile Asn  Ala Trp Ser Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val
    1460                1465                1470

Asp Leu  Leu Ala Pro Met Ile  Ile His Gly Ile Lys  Thr Gln Gly
    1475                1480                1485

Ala Arg  Gln Lys Phe Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile
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Met Tyr  Ser Leu Asp Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn
    1505                1510                1515

Ser Thr  Gly Thr Leu Met Val  Phe Phe Gly Asn Val  Asp Ser Ser
    1520                1525                1530

Gly Ile  Lys His Asn Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr
    1535                1540                1545

Ile Arg  Leu His Pro Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg
    1550                1555                1560

Met Glu  Leu Met Gly Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu
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Gly Met  Glu Ser Lys Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser
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Ser Tyr  Phe Thr Asn Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala
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Arg Leu  His Leu Gln Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val
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Asn Asn  Pro Lys Glu Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met
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Lys Val  Thr Gly Val Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr
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Ser Met  Tyr Val Lys Glu Phe  Leu Ile Ser Ser Ser  Gln Asp Gly
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His Gln  Trp Thr Leu Phe Phe  Gln Asn Gly Lys Val  Lys Val Phe
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Gln Gly  Asn Gln Asp Ser Phe  Thr Pro Val Val Asn  Ser Leu Asp
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Pro Pro  Leu Leu Thr Arg Tyr  Leu Arg Ile His Pro  Gln Ser Trp
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Val His  Gln Ile Ala Leu Arg  Met Glu Val Leu Gly  Cys Glu Ala
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Gln Asp  Leu Tyr Gly Ala Pro  Thr Ser Thr Glu Pro  Ser Glu Gly
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Ser Ala  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
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Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
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Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
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Ser Gly  Ser Glu Thr Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
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Glu Thr  Pro Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
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Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1820                1825                1830

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Glu  Pro Ala Thr
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Ser Gly  Ser Glu Thr Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
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Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
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Gly Ala  Ser Ser Glu Gln Lys  Leu Ile Ser Glu Glu  Asp Leu Ser
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

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Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495
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His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Pro Gly Ala Ser
            740                 745                 750

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
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Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
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Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro
785                 790                 795                 800

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
                805                 810                 815

Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
            820                 825                 830

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
        835                 840                 845

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
    850                 855                 860

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
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Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
                885                 890                 895

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
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Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
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Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
945                 950                 955                 960

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
            980                 985                 990

Pro Gly Thr Ser Ser Thr Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser
        995                 1000                1005

Thr Gly  Thr Gly Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
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Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Ala Ser
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Ser Pro  Pro Val Leu Lys Arg  His Gln Arg Glu Ile  Thr Arg Thr
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Thr Leu  Gln Ser Asp Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile
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Ser Val  Glu Met Lys Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp
    1070                1075                1080

Glu Asn  Gln Ser Pro Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr
    1085                1090                1095

Phe Ile  Ala Ala Val Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser
    1100                1105                1110

Ser Pro  His Val Leu Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro
    1115                1120                1125

Gln Phe  Lys Lys Val Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe
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Thr Gln  Pro Leu Tyr Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu
    1145                1150                1155

Leu Gly  Pro Tyr Ile Arg Ala  Glu Val Glu Asp Asn  Ile Met Val
    1160                1165                1170

Thr Phe  Arg Asn Gln Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser
    1175                1180                1185

Leu Ile  Ser Tyr Glu Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg
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Lys Asn  Phe Val Lys Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys
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Val Gln  His His Met Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys
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Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu Glu Lys  Asp Val His
    1235                1240                1245

Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys His Thr  Asn Thr Leu
    1250                1255                1260

Asn Pro  Ala His Gly Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu
    1265                1270                1275

Phe Phe  Thr Ile Phe Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu
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Asn Met  Glu Arg Asn Cys Arg  Ala Pro Cys Asn Ile  Gln Met Glu
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Asp Pro  Thr Phe Lys Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly
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Tyr Ile  Met Asp Thr Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln
    1325                1330                1335

Arg Ile  Arg Trp Tyr Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile
    1340                1345                1350

His Ser  Ile His Phe Ser Gly  His Val Phe Thr Val  Arg Lys Lys
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Glu Glu  Tyr Lys Met Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe
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Glu Thr  Val Glu Met Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val
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Glu Cys  Leu Ile Gly Glu His  Leu His Ala Gly Met  Ser Thr Leu
    1400                1405                1410

Phe Leu  Val Tyr Ser Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala
    1415                1420                1425

Ser Gly  His Ile Arg Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr
    1430                1435                1440

Gly Gln  Trp Ala Pro Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser
    1445                1450                1455

Ile Asn  Ala Trp Ser Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val
    1460                1465                1470

Asp Leu  Leu Ala Pro Met Ile  Ile His Gly Ile Lys  Thr Gln Gly
    1475                1480                1485

Ala Arg  Gln Lys Phe Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile
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Met Tyr  Ser Leu Asp Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn
    1505                1510                1515

Ser Thr  Gly Thr Leu Met Val  Phe Phe Gly Asn Val  Asp Ser Ser
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Gly Ile  Lys His Asn Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr
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Ile Arg  Leu His Pro Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg
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Met Glu  Leu Met Gly Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu
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Gly Met  Glu Ser Lys Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser
    1580                1585                1590

Ser Tyr  Phe Thr Asn Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala
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Arg Leu  His Leu Gln Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val
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Asn Asn  Pro Lys Glu Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met
    1625                1630                1635

Lys Val  Thr Gly Val Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr
    1640                1645                1650

Ser Met  Tyr Val Lys Glu Phe  Leu Ile Ser Ser Ser  Gln Asp Gly
    1655                1660                1665

His Gln  Trp Thr Leu Phe Phe  Gln Asn Gly Lys Val  Lys Val Phe
    1670                1675                1680

Gln Gly  Asn Gln Asp Ser Phe  Thr Pro Val Val Asn  Ser Leu Asp
    1685                1690                1695

Pro Pro  Leu Leu Thr Arg Tyr  Leu Arg Ile His Pro  Gln Ser Trp
    1700                1705                1710

Val His  Gln Ile Ala Leu Arg  Met Glu Val Leu Gly  Cys Glu Ala
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Gln Asp  Leu Tyr Gly Ala Pro  Thr Ser Thr Glu Pro  Ser Glu Gly
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Ser Ala  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
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Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
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Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
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Ser Gly  Ser Glu Thr Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1790                1795                1800
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Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
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Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
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Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
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<212> PRT
<213> Artificial Sequence

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20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val

115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu

385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro

660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Thr Ser Glu
740 745 750

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
755 760 765

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770 775 780

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785 790 795 800

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805 810 815

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820 825 830

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
835 840 845

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
850 855 860

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865 870 875 880

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885 890 895

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
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Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
915 920 925

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro

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            980                 985                 990

Glu Pro Ser Glu Gly Ser Ala Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
        995                 1000                1005

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
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Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Ala Ser
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Ser Pro  Pro Val Leu Lys Arg  His Gln Arg Glu Ile  Thr Arg Thr
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Thr Leu  Gln Ser Asp Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile
    1055                1060                1065

Ser Val  Glu Met Lys Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp
    1070                1075                1080

Glu Asn  Gln Ser Pro Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr
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Phe Ile  Ala Ala Val Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser
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Ser Pro  His Val Leu Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro
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Gln Phe  Lys Lys Val Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe
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Thr Gln  Pro Leu Tyr Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu
    1145                1150                1155

Leu Gly  Pro Tyr Ile Arg Ala  Glu Val Glu Asp Asn  Ile Met Val
    1160                1165                1170

Thr Phe  Arg Asn Gln Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser
    1175                1180                1185

Leu Ile  Ser Tyr Glu Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg
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Val Gln His His Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys
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Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu Lys Asp Val His
1235 1240 1245

Ser Gly Leu Ile Gly Pro Leu Leu Val Cys His Thr Asn Thr Leu
1250 1255 1260

Asn Pro Ala His Gly Arg Gln Val Thr Val Gln Glu Phe Ala Leu
1265 1270 1275

Phe Phe Thr Ile Phe Asp Glu Thr Lys Ser Trp Tyr Phe Thr Glu
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Asn Met Glu Arg Asn Cys Arg Ala Pro Cys Asn Ile Gln Met Glu
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Asp Pro Thr Phe Lys Glu Asn Tyr Arg Phe His Ala Ile Asn Gly
1310 1315 1320

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Arg Ile Arg Trp Tyr Leu Leu Ser Met Gly Ser Asn Glu Asn Ile
1340 1345 1350

His Ser Ile His Phe Ser Gly His Val Phe Thr Val Arg Lys Lys
1355 1360 1365

Glu Glu Tyr Lys Met Ala Leu Tyr Asn Leu Tyr Pro Gly Val Phe
1370 1375 1380

Glu Thr Val Glu Met Leu Pro Ser Lys Ala Gly Ile Trp Arg Val
1385 1390 1395

Glu Cys Leu Ile Gly Glu His Leu His Ala Gly Met Ser Thr Leu
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Phe Leu Val Tyr Ser Asn Lys Cys Gln Thr Pro Leu Gly Met Ala
1415 1420 1425

Ser Gly His Ile Arg Asp Phe Gln Ile Thr Ala Ser Gly Gln Tyr
1430 1435 1440

Gly Gln Trp Ala Pro Lys Leu Ala Arg Leu His Tyr Ser Gly Ser

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Ile Asn  Ala Trp Ser Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val
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Asp Leu  Leu Ala Pro Met Ile  Ile His Gly Ile Lys  Thr Gln Gly
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Ala Arg  Gln Lys Phe Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile
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Met Tyr  Ser Leu Asp Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn
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Ser Thr  Gly Thr Leu Met Val  Phe Phe Gly Asn Val  Asp Ser Ser
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Gly Ile  Lys His Asn Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr
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Ile Arg  Leu His Pro Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg
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Met Glu  Leu Met Gly Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu
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Gly Met  Glu Ser Lys Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser
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Ser Tyr  Phe Thr Asn Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala
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Arg Leu  His Leu Gln Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val
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Asn Asn  Pro Lys Glu Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met
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Lys Val  Thr Gly Val Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr
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Ser Met  Tyr Val Lys Glu Phe  Leu Ile Ser Ser Ser  Gln Asp Gly
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His Gln  Trp Thr Leu Phe Phe  Gln Asn Gly Lys Val  Lys Val Phe
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Gln Gly  Asn Gln Asp Ser Phe  Thr Pro Val Val Asn  Ser Leu Asp
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Pro Pro  Leu Leu Thr Arg Tyr  Leu Arg Ile His Pro  Gln Ser Trp
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Gln Asp Leu Tyr Gly Ala Pro Pro Gly Ser Ser Pro Ser Ala Ser
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Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr
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1805 1810 1815

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Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
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Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
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Pro Ala Thr Gly His His His His His His
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<210> 827
<211> 1903
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285
```

```
Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560
```

```
Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Pro Gly Ala Ser
            740                 745                 750

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
        755                 760                 765

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
    770                 775                 780

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro
785                 790                 795                 800

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
                805                 810                 815

Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
            820                 825                 830
```

```
Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
        835                 840                 845

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
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Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
865                 870                 875                 880

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
                885                 890                 895

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
            900                 905                 910

Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
        915                 920                 925

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
    930                 935                 940

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
945                 950                 955                 960

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
                965                 970                 975

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
            980                 985                 990

Pro Gly Thr Ser Ser Thr Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser
        995                 1000                1005

Thr Gly  Thr Gly Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
    1010                1015                1020

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Ala Ser
    1025                1030                1035

Ser Pro  Pro Val Leu Lys Arg  His Gln Arg Glu Ile  Thr Arg Thr
    1040                1045                1050

Thr Leu  Gln Ser Asp Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile
    1055                1060                1065

Ser Val  Glu Met Lys Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp
    1070                1075                1080

Glu Asn  Gln Ser Pro Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr
    1085                1090                1095
```

```
Phe Ile  Ala Ala Val Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser
    1100                1105                1110

Ser Pro  His Val Leu Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro
    1115                1120                1125

Gln Phe  Lys Lys Val Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe
    1130                1135                1140

Thr Gln  Pro Leu Tyr Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu
    1145                1150                1155

Leu Gly  Pro Tyr Ile Arg Ala  Glu Val Glu Asp Asn  Ile Met Val
    1160                1165                1170

Thr Phe  Arg Asn Gln Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser
    1175                1180                1185

Leu Ile  Ser Tyr Glu Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg
    1190                1195                1200

Lys Asn  Phe Val Lys Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys
    1205                1210                1215

Val Gln  His His Met Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys
    1220                1225                1230

Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu Glu Lys  Asp Val His
    1235                1240                1245

Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys His Thr  Asn Thr Leu
    1250                1255                1260

Asn Pro  Ala His Gly Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu
    1265                1270                1275

Phe Phe  Thr Ile Phe Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu
    1280                1285                1290

Asn Met  Glu Arg Asn Cys Arg  Ala Pro Cys Asn Ile  Gln Met Glu
    1295                1300                1305

Asp Pro  Thr Phe Lys Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly
    1310                1315                1320

Tyr Ile  Met Asp Thr Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln
    1325                1330                1335

Arg Ile  Arg Trp Tyr Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile
    1340                1345                1350
```

```
His Ser  Ile His Phe Ser Gly  His Val Phe Thr Val  Arg Lys Lys
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Glu Glu  Tyr Lys Met Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe
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Glu Thr  Val Glu Met Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val
    1385                1390                1395

Glu Cys  Leu Ile Gly Glu His  Leu His Ala Gly Met  Ser Thr Leu
    1400                1405                1410

Phe Leu  Val Tyr Ser Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala
    1415                1420                1425

Ser Gly  His Ile Arg Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr
    1430                1435                1440

Gly Gln  Trp Ala Pro Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser
    1445                1450                1455

Ile Asn  Ala Trp Ser Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val
    1460                1465                1470

Asp Leu  Leu Ala Pro Met Ile  Ile His Gly Ile Lys  Thr Gln Gly
    1475                1480                1485

Ala Arg  Gln Lys Phe Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile
    1490                1495                1500

Met Tyr  Ser Leu Asp Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn
    1505                1510                1515

Ser Thr  Gly Thr Leu Met Val  Phe Phe Gly Asn Val  Asp Ser Ser
    1520                1525                1530

Gly Ile  Lys His Asn Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr
    1535                1540                1545

Ile Arg  Leu His Pro Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg
    1550                1555                1560

Met Glu  Leu Met Gly Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu
    1565                1570                1575

Gly Met  Glu Ser Lys Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser
    1580                1585                1590

Ser Tyr  Phe Thr Asn Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala
    1595                1600                1605
```

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Arg Leu  His Leu Gln Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val
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Asn Asn  Pro Lys Glu Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met
    1625                 1630                 1635

Lys Val  Thr Gly Val Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr
    1640                 1645                 1650

Ser Met  Tyr Val Lys Glu Phe  Leu Ile Ser Ser Ser  Gln Asp Gly
    1655                 1660                 1665

His Gln  Trp Thr Leu Phe Phe  Gln Asn Gly Lys Val  Lys Val Phe
    1670                 1675                 1680

Gln Gly  Asn Gln Asp Ser Phe  Thr Pro Val Val Asn  Ser Leu Asp
    1685                 1690                 1695

Pro Pro  Leu Leu Thr Arg Tyr  Leu Arg Ile His Pro  Gln Ser Trp
    1700                 1705                 1710

Val His  Gln Ile Ala Leu Arg  Met Glu Val Leu Gly  Cys Glu Ala
    1715                 1720                 1725

Gln Asp  Leu Tyr Gly Ala Pro  Pro Gly Ser Ser Pro  Ser Ala Ser
    1730                 1735                 1740

Thr Gly  Thr Gly Pro Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr
    1745                 1750                 1755

Gly Pro  Gly Thr Pro Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly
    1760                 1765                 1770

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Pro
    1775                 1780                 1785

Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ala Ser Pro  Gly Thr Ser
    1790                 1795                 1800

Ser Thr  Gly Ser Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
    1805                 1810                 1815

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly
    1820                 1825                 1830

Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser Pro Gly  Ala Ser Pro
    1835                 1840                 1845

Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1850                 1855                 1860
```

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
1865 1870 1875

Ser Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Ala Thr Gly His His His His His His
1895 1900

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acgggttcgc cgggaagcag cccaagcgca tccactggta cagggcctgg ggcttcaccg 5400

ggtacttcat ccacggggtc accgggaacg cccggatcgg ggacggcttc ctcatcacca 5460

ggatcgtcaa caccctcggg cgcaacgggc agccccggaa cccctggttc gggtacggcg 5520

tcgtcgagcc ccggtgcgag cccgggaaca agctcgacag gatcgcctgg ggcgtcaccc 5580

ggcacgtcga gcacaggcag ccccggaacc cctggatcgg gaaccgcgtc gtcaagcgcc 5640

tcgagcgaac aaaaactcat ctcagaagag gatctgtcac ctgcaaccgg tcatcatcac 5700

catcaccatt ga 5712

<210> 829
<211> 1903
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 829

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

```
His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460
```

```
Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735
```

```
Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Thr Ser Glu
            740                 745                 750

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
        755                 760                 765

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
    770                 775                 780

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
785                 790                 795                 800

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
                805                 810                 815

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
            820                 825                 830

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
        835                 840                 845

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
    850                 855                 860

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
865                 870                 875                 880

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
                885                 890                 895

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
            900                 905                 910

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        915                 920                 925

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
    930                 935                 940

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
945                 950                 955                 960

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
                965                 970                 975

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
            980                 985                 990

Glu Pro Ser Glu Gly Ser Ala Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
        995                 1000                1005
```

```
Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1010                1015                1020

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Ala Ser
    1025                1030                1035

Ser Pro  Pro Val Leu Lys Arg  His Gln Ala Glu Ile  Thr Arg Thr
    1040                1045                1050

Thr Leu  Gln Ser Asp Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile
    1055                1060                1065

Ser Val  Glu Met Lys Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp
    1070                1075                1080

Glu Asn  Gln Ser Pro Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr
    1085                1090                1095

Phe Ile  Ala Ala Val Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser
    1100                1105                1110

Ser Pro  His Val Leu Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro
    1115                1120                1125

Gln Phe  Lys Lys Val Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe
    1130                1135                1140

Thr Gln  Pro Leu Tyr Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu
    1145                1150                1155

Leu Gly  Pro Tyr Ile Arg Ala  Glu Val Glu Asp Asn  Ile Met Val
    1160                1165                1170

Thr Phe  Arg Asn Gln Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser
    1175                1180                1185

Leu Ile  Ser Tyr Glu Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg
    1190                1195                1200

Lys Asn  Phe Val Lys Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys
    1205                1210                1215

Val Gln  His His Met Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys
    1220                1225                1230

Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu Glu Lys  Asp Val His
    1235                1240                1245

Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys His Thr  Asn Thr Leu
    1250                1255                1260
```

```
Asn Pro  Ala His Gly Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu
    1265                 1270                 1275

Phe Phe  Thr Ile Phe Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu
    1280                 1285                 1290

Asn Met  Glu Arg Asn Cys Arg  Ala Pro Cys Asn Ile  Gln Met Glu
    1295                 1300                 1305

Asp Pro  Thr Phe Lys Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly
    1310                 1315                 1320

Tyr Ile  Met Asp Thr Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln
    1325                 1330                 1335

Arg Ile  Arg Trp Tyr Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile
    1340                 1345                 1350

His Ser  Ile His Phe Ser Gly  His Val Phe Thr Val  Arg Lys Lys
    1355                 1360                 1365

Glu Glu  Tyr Lys Met Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe
    1370                 1375                 1380

Glu Thr  Val Glu Met Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val
    1385                 1390                 1395

Glu Cys  Leu Ile Gly Glu His  Leu His Ala Gly Met  Ser Thr Leu
    1400                 1405                 1410

Phe Leu  Val Tyr Ser Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala
    1415                 1420                 1425

Ser Gly  His Ile Arg Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr
    1430                 1435                 1440

Gly Gln  Trp Ala Pro Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser
    1445                 1450                 1455

Ile Asn  Ala Trp Ser Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val
    1460                 1465                 1470

Asp Leu  Leu Ala Pro Met Ile  Ile His Gly Ile Lys  Thr Gln Gly
    1475                 1480                 1485

Ala Arg  Gln Lys Phe Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile
    1490                 1495                 1500

Met Tyr  Ser Leu Asp Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn
    1505                 1510                 1515
```

```
Ser Thr  Gly Thr Leu Met Val  Phe Phe Gly Asn Val  Asp Ser Ser
    1520                1525                1530

Gly Ile  Lys His Asn Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr
    1535                1540                1545

Ile Arg  Leu His Pro Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg
    1550                1555                1560

Met Glu  Leu Met Gly Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu
    1565                1570                1575

Gly Met  Glu Ser Lys Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser
    1580                1585                1590

Ser Tyr  Phe Thr Asn Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala
    1595                1600                1605

Arg Leu  His Leu Gln Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val
    1610                1615                1620

Asn Asn  Pro Lys Glu Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met
    1625                1630                1635

Lys Val  Thr Gly Val Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr
    1640                1645                1650

Ser Met  Tyr Val Lys Glu Phe  Leu Ile Ser Ser Ser  Gln Asp Gly
    1655                1660                1665

His Gln  Trp Thr Leu Phe Phe  Gln Asn Gly Lys Val  Lys Val Phe
    1670                1675                1680

Gln Gly  Asn Gln Asp Ser Phe  Thr Pro Val Val Asn  Ser Leu Asp
    1685                1690                1695

Pro Pro  Leu Leu Thr Arg Tyr  Leu Arg Ile His Pro  Gln Ser Trp
    1700                1705                1710

Val His  Gln Ile Ala Leu Arg  Met Glu Val Leu Gly  Cys Glu Ala
    1715                1720                1725

Gln Asp  Leu Tyr Gly Ala Pro  Pro Gly Ser Ser Pro  Ser Ala Ser
    1730                1735                1740

Thr Gly  Thr Gly Pro Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr
    1745                1750                1755

Gly Pro  Gly Thr Pro Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly
    1760                1765                1770
```

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
1775 1780 1785

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser
1790 1795 1800

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
1805 1810 1815

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
1820 1825 1830

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
1835 1840 1845

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
1850 1855 1860

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
1865 1870 1875

Ser Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
1880 1885 1890

Pro Ala Thr Gly His His His His His His
1895 1900

<210> 830
<211> 5712
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

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gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

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catcaccatt ga 5712

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<211> 2047
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

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Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365
```

```
Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640
```

```
Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Thr Ser Glu
            740                 745                 750

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
        755                 760                 765

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
    770                 775                 780

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
785                 790                 795                 800

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
                805                 810                 815

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
            820                 825                 830

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
        835                 840                 845

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
    850                 855                 860

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
865                 870                 875                 880

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
                885                 890                 895

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
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Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
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Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
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Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
945                 950                 955                 960

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
                965                 970                 975

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
            980                 985                 990

Glu Pro Ser Glu Gly Ser Ala Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
        995                 1000                1005

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1010                1015                1020

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Ala Ser
    1025                1030                1035

Ser Pro  Pro Val Leu Lys Arg  His Gln Ala Glu Ile  Thr Arg Thr
    1040                1045                1050

Thr Leu  Gln Ser Asp Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile
    1055                1060                1065

Ser Val  Glu Met Lys Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp
    1070                1075                1080

Glu Asn  Gln Ser Pro Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr
    1085                1090                1095

Phe Ile  Ala Ala Val Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser
    1100                1105                1110

Ser Pro  His Val Leu Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro
    1115                1120                1125

Gln Phe  Lys Lys Val Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe
    1130                1135                1140

Thr Gln  Pro Leu Tyr Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu
    1145                1150                1155

Leu Gly  Pro Tyr Ile Arg Ala  Glu Val Glu Asp Asn  Ile Met Val
    1160                1165                1170
```

```
Thr Phe  Arg Asn Gln Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser
    1175                1180                1185

Leu Ile  Ser Tyr Glu Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg
    1190                1195                1200

Lys Asn  Phe Val Lys Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys
    1205                1210                1215

Val Gln  His His Met Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys
    1220                1225                1230

Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu Glu Lys  Asp Val His
    1235                1240                1245

Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys His Thr  Asn Thr Leu
    1250                1255                1260

Asn Pro  Ala His Gly Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu
    1265                1270                1275

Phe Phe  Thr Ile Phe Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu
    1280                1285                1290

Asn Met  Glu Arg Asn Cys Arg  Ala Pro Cys Asn Ile  Gln Met Glu
    1295                1300                1305

Asp Pro  Thr Phe Lys Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly
    1310                1315                1320

Tyr Ile  Met Asp Thr Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln
    1325                1330                1335

Arg Ile  Arg Trp Tyr Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile
    1340                1345                1350

His Ser  Ile His Phe Ser Gly  His Val Phe Thr Val  Arg Lys Lys
    1355                1360                1365

Glu Glu  Tyr Lys Met Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe
    1370                1375                1380

Glu Thr  Val Glu Met Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val
    1385                1390                1395

Glu Cys  Leu Ile Gly Glu His  Leu His Ala Gly Met  Ser Thr Leu
    1400                1405                1410

Phe Leu  Val Tyr Ser Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala
    1415                1420                1425
```

```
Ser Gly  His Ile Arg Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr
    1430                1435                1440

Gly Gln  Trp Ala Pro Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser
    1445                1450                1455

Ile Asn  Ala Trp Ser Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val
    1460                1465                1470

Asp Leu  Leu Ala Pro Met Ile  Ile His Gly Ile Lys  Thr Gln Gly
    1475                1480                1485

Ala Arg  Gln Lys Phe Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile
    1490                1495                1500

Met Tyr  Ser Leu Asp Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn
    1505                1510                1515

Ser Thr  Gly Thr Leu Met Val  Phe Phe Gly Asn Val  Asp Ser Ser
    1520                1525                1530

Gly Ile  Lys His Asn Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr
    1535                1540                1545

Ile Arg  Leu His Pro Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg
    1550                1555                1560

Met Glu  Leu Met Gly Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu
    1565                1570                1575

Gly Met  Glu Ser Lys Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser
    1580                1585                1590

Ser Tyr  Phe Thr Asn Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala
    1595                1600                1605

Arg Leu  His Leu Gln Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val
    1610                1615                1620

Asn Asn  Pro Lys Glu Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met
    1625                1630                1635

Lys Val  Thr Gly Val Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr
    1640                1645                1650

Ser Met  Tyr Val Lys Glu Phe  Leu Ile Ser Ser Ser  Gln Asp Gly
    1655                1660                1665

His Gln  Trp Thr Leu Phe Phe  Gln Asn Gly Lys Val  Lys Val Phe
    1670                1675                1680
```

```
Gln Gly  Asn Gln Asp Ser Phe  Thr Pro Val Val Asn  Ser Leu Asp
    1685                1690                1695

Pro Pro  Leu Leu Thr Arg Tyr  Leu Arg Ile His Pro  Gln Ser Trp
    1700                1705                1710

Val His  Gln Ile Ala Leu Arg  Met Glu Val Leu Gly  Cys Glu Ala
    1715                1720                1725

Gln Asp  Leu Tyr Gly Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1730                1735                1740

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1745                1750                1755

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1760                1765                1770

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    1775                1780                1785

Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu
    1790                1795                1800

Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    1805                1810                1815

Glu Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1820                1825                1830

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    1835                1840                1845

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Pro Ala Gly  Ser Pro Thr
    1850                1855                1860

Ser Thr  Glu Glu Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    1865                1870                1875

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Thr  Ser Glu Ser
    1895                1900                1905

Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935
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Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser
1940 1945 1950

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu
1955 1960 1965

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
1970 1975 1980

Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1985 1990 1995

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
2000 2005 2010

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Ala Ser Ser Glu Gln
2015 2020 2025

Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala Thr Gly His His
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His His His His
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20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

```
Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320
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Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
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Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
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Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590
```

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Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
            740                 745                 750

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
        755                 760                 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
    770                 775                 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785                 790                 795                 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                805                 810                 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            820                 825                 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
        835                 840                 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    850                 855                 860
```

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Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865                 870                 875                 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125
```

```
Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
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Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380
```

```
Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                1435                1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                1450                1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                1465                1470

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                1480                1485

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1490                1495                1500

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                1510                1515

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1520                1525                1530

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1535                1540                1545

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1550                1555                1560

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1565                1570                1575

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1580                1585                1590

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1595                1600                1605

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1610                1615                1620

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1625                1630                1635
```

```
Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1640                 1645                 1650

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1655                 1660                 1665

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1670                 1675                 1680

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1685                 1690                 1695

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1700                 1705                 1710

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    1715                 1720                 1725

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    1730                 1735                 1740

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1745                 1750                 1755

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1760                 1765                 1770

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1775                 1780                 1785

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1790                 1795                 1800

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    1805                 1810                 1815

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    1820                 1825                 1830

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Glu Pro Ala
    1835                 1840                 1845

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1850                 1855                 1860

Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    1865                 1870                 1875

Pro Ala  Ser Ser Glu Gln Lys  Leu Ile Ser Glu Glu  Asp Leu Ser
    1880                 1885                 1890
```

Pro Ala Thr Gly His His His His His His
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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

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<211> 1903
<212> PRT
<213> Artificial Sequence

<220>
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20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

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Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495
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His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
            740                 745                 750

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
        755                 760                 765
```

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Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
    770                 775                 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785                 790                 795                 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                805                 810                 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            820                 825                 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
        835                 840                 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    850                 855                 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865                 870                 875                 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035
```

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1040 1045 1050

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1055 1060 1065

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1070 1075 1080

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1085 1090 1095

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1100 1105 1110

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1115 1120 1125

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1130 1135 1140

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1145 1150 1155

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1160 1165 1170

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1175 1180 1185

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1190 1195 1200

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1205 1210 1215

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1220 1225 1230

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1235 1240 1245

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1250 1255 1260

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1265 1270 1275

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1280 1285 1290

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Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                1435                1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                1450                1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                1465                1470

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                1480                1485

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1490                1495                1500

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                1510                1515

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1520                1525                1530

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1535                1540                1545
```

```
Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1550                1555                1560

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1565                1570                1575

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1580                1585                1590

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1595                1600                1605

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1610                1615                1620

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1625                1630                1635

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1640                1645                1650

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1655                1660                1665

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1670                1675                1680

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1685                1690                1695

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1700                1705                1710

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    1715                1720                1725

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    1730                1735                1740

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1745                1750                1755

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1760                1765                1770

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1775                1780                1785

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1790                1795                1800
```

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
1805 1810 1815

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
1820 1825 1830

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
1835 1840 1845

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
1850 1855 1860

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
1865 1870 1875

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
1880 1885 1890

Pro Ala Thr Gly His His His His His His
1895 1900

<210> 836
<211> 5712
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 836
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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

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gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

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actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

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tcagcacccg gctctgagcc ggccacaagt ggcagtgaga cacccggcac ttcagagagt 5580

gccacccccg agagtggccc aggcactagt accgagccct ctgaaggcag tgcgccagcc 5640

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catcaccatt ga 5712

<210> 837
<211> 2047
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 837

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val

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        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
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385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro

660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Pro Gly Ala Ser
740 745 750

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
755 760 765

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
770 775 780

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro
785 790 795 800

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
805 810 815

Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
820 825 830

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
835 840 845

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
850 855 860

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
865 870 875 880

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
885 890 895

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
900 905 910

Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
915 920 925

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser

930 935 940

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
945 950 955 960

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
965 970 975

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
980 985 990

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
995 1000 1005

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
1010 1015 1020

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Ala Ser
1025 1030 1035

Ser Pro Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr
1040 1045 1050

Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
1055 1060 1065

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp
1070 1075 1080

Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr
1085 1090 1095

Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser
1100 1105 1110

Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro
1115 1120 1125

Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe
1130 1135 1140

Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
1145 1150 1155

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val
1160 1165 1170

Thr Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser
1175 1180 1185

Leu Ile Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg

1190 1195 1200

Lys Asn Phe Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys
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Val Gln His His Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys
1220 1225 1230

Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu Lys Asp Val His
1235 1240 1245

Ser Gly Leu Ile Gly Pro Leu Leu Val Cys His Thr Asn Thr Leu
1250 1255 1260

Asn Pro Ala His Gly Arg Gln Val Thr Val Gln Glu Phe Ala Leu
1265 1270 1275

Phe Phe Thr Ile Phe Asp Glu Thr Lys Ser Trp Tyr Phe Thr Glu
1280 1285 1290

Asn Met Glu Arg Asn Cys Arg Ala Pro Cys Asn Ile Gln Met Glu
1295 1300 1305

Asp Pro Thr Phe Lys Glu Asn Tyr Arg Phe His Ala Ile Asn Gly
1310 1315 1320

Tyr Ile Met Asp Thr Leu Pro Gly Leu Val Met Ala Gln Asp Gln
1325 1330 1335

Arg Ile Arg Trp Tyr Leu Leu Ser Met Gly Ser Asn Glu Asn Ile
1340 1345 1350

His Ser Ile His Phe Ser Gly His Val Phe Thr Val Arg Lys Lys
1355 1360 1365

Glu Glu Tyr Lys Met Ala Leu Tyr Asn Leu Tyr Pro Gly Val Phe
1370 1375 1380

Glu Thr Val Glu Met Leu Pro Ser Lys Ala Gly Ile Trp Arg Val
1385 1390 1395

Glu Cys Leu Ile Gly Glu His Leu His Ala Gly Met Ser Thr Leu
1400 1405 1410

Phe Leu Val Tyr Ser Asn Lys Cys Gln Thr Pro Leu Gly Met Ala
1415 1420 1425

Ser Gly His Ile Arg Asp Phe Gln Ile Thr Ala Ser Gly Gln Tyr
1430 1435 1440

Gly Gln Trp Ala Pro Lys Leu Ala Arg Leu His Tyr Ser Gly Ser

1445 1450 1455

Ile Asn Ala Trp Ser Thr Lys Glu Pro Phe Ser Trp Ile Lys Val
1460 1465 1470

Asp Leu Leu Ala Pro Met Ile Ile His Gly Ile Lys Thr Gln Gly
1475 1480 1485

Ala Arg Gln Lys Phe Ser Ser Leu Tyr Ile Ser Gln Phe Ile Ile
1490 1495 1500

Met Tyr Ser Leu Asp Gly Lys Lys Trp Gln Thr Tyr Arg Gly Asn
1505 1510 1515

Ser Thr Gly Thr Leu Met Val Phe Phe Gly Asn Val Asp Ser Ser
1520 1525 1530

Gly Ile Lys His Asn Ile Phe Asn Pro Pro Ile Ile Ala Arg Tyr
1535 1540 1545

Ile Arg Leu His Pro Thr His Tyr Ser Ile Arg Ser Thr Leu Arg
1550 1555 1560

Met Glu Leu Met Gly Cys Asp Leu Asn Ser Cys Ser Met Pro Leu
1565 1570 1575

Gly Met Glu Ser Lys Ala Ile Ser Asp Ala Gln Ile Thr Ala Ser
1580 1585 1590

Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp Ser Pro Ser Lys Ala
1595 1600 1605

Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg Pro Gln Val
1610 1615 1620

Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln Lys Thr Met
1625 1630 1635

Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser Leu Leu Thr
1640 1645 1650

Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly
1655 1660 1665

His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val Lys Val Phe
1670 1675 1680

Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn Ser Leu Asp
1685 1690 1695

Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro Gln Ser Trp

1700 1705 1710

Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly Cys Glu Ala
1715 1720 1725

Gln Asp Leu Tyr Gly Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
1730 1735 1740

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1745 1750 1755

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
1760 1765 1770

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
1775 1780 1785

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
1790 1795 1800

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
1805 1810 1815

Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
1820 1825 1830

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
1835 1840 1845

Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
1850 1855 1860

Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
1865 1870 1875

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
1880 1885 1890

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser
1895 1900 1905

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
1910 1915 1920

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1925 1930 1935

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser
1940 1945 1950

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu

1955 1960 1965

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
1970 1975 1980

Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1985 1990 1995

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
2000 2005 2010

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Ala Ser Ser Glu Gln
2015 2020 2025

Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala Thr Gly His His
2030 2035 2040

His His His His
2045

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<211> 6144
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<213> Artificial Sequence

<220>
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atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

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tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

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gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

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ggtcatcatc accatcacca ttga 6144

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<211> 2047
<212> PRT
<213> Artificial Sequence

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

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Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350
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Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620
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Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Thr Ser Glu
            740                 745                 750

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
        755                 760                 765

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
    770                 775                 780

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
785                 790                 795                 800

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
                805                 810                 815

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
            820                 825                 830

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
        835                 840                 845

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
    850                 855                 860

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
865                 870                 875                 880

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
                885                 890                 895
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Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
            900                 905                 910

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        915                 920                 925

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
    930                 935                 940

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
945                 950                 955                 960

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
                965                 970                 975

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
            980                 985                 990

Glu Pro Ser Glu Gly Ser Ala Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
        995                 1000                1005

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1010                1015                1020

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Ala Ser
    1025                1030                1035

Ser Pro  Pro Val Leu Lys Arg  His Gln Arg Glu Ile  Thr Arg Thr
    1040                1045                1050

Thr Leu  Gln Ser Asp Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile
    1055                1060                1065

Ser Val  Glu Met Lys Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp
    1070                1075                1080

Glu Asn  Gln Ser Pro Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr
    1085                1090                1095

Phe Ile  Ala Ala Val Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser
    1100                1105                1110

Ser Pro  His Val Leu Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro
    1115                1120                1125

Gln Phe  Lys Lys Val Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe
    1130                1135                1140

Thr Gln  Pro Leu Tyr Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu
    1145                1150                1155
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Leu Gly  Pro Tyr Ile Arg Ala  Glu Val Glu Asp Asn  Ile Met Val
    1160                1165                1170

Thr Phe  Arg Asn Gln Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser
    1175                1180                1185

Leu Ile  Ser Tyr Glu Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg
    1190                1195                1200

Lys Asn  Phe Val Lys Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys
    1205                1210                1215

Val Gln  His His Met Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys
    1220                1225                1230

Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu Glu Lys  Asp Val His
    1235                1240                1245

Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys His Thr  Asn Thr Leu
    1250                1255                1260

Asn Pro  Ala His Gly Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu
    1265                1270                1275

Phe Phe  Thr Ile Phe Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu
    1280                1285                1290

Asn Met  Glu Arg Asn Cys Arg  Ala Pro Cys Asn Ile  Gln Met Glu
    1295                1300                1305

Asp Pro  Thr Phe Lys Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly
    1310                1315                1320

Tyr Ile  Met Asp Thr Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln
    1325                1330                1335

Arg Ile  Arg Trp Tyr Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile
    1340                1345                1350

His Ser  Ile His Phe Ser Gly  His Val Phe Thr Val  Arg Lys Lys
    1355                1360                1365

Glu Glu  Tyr Lys Met Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe
    1370                1375                1380

Glu Thr  Val Glu Met Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val
    1385                1390                1395

Glu Cys  Leu Ile Gly Glu His  Leu His Ala Gly Met  Ser Thr Leu
    1400                1405                1410
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Phe Leu  Val Tyr Ser Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala
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Ser Gly  His Ile Arg Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr
    1430                1435                1440

Gly Gln  Trp Ala Pro Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser
    1445                1450                1455

Ile Asn  Ala Trp Ser Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val
    1460                1465                1470

Asp Leu  Leu Ala Pro Met Ile  Ile His Gly Ile Lys  Thr Gln Gly
    1475                1480                1485

Ala Arg  Gln Lys Phe Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile
    1490                1495                1500

Met Tyr  Ser Leu Asp Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn
    1505                1510                1515

Ser Thr  Gly Thr Leu Met Val  Phe Phe Gly Asn Val  Asp Ser Ser
    1520                1525                1530

Gly Ile  Lys His Asn Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr
    1535                1540                1545

Ile Arg  Leu His Pro Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg
    1550                1555                1560

Met Glu  Leu Met Gly Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu
    1565                1570                1575

Gly Met  Glu Ser Lys Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser
    1580                1585                1590

Ser Tyr  Phe Thr Asn Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala
    1595                1600                1605

Arg Leu  His Leu Gln Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val
    1610                1615                1620

Asn Asn  Pro Lys Glu Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met
    1625                1630                1635

Lys Val  Thr Gly Val Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr
    1640                1645                1650

Ser Met  Tyr Val Lys Glu Phe  Leu Ile Ser Ser Ser  Gln Asp Gly
    1655                1660                1665
```

```
His Gln  Trp Thr Leu Phe Phe  Gln Asn Gly Lys Val  Lys Val Phe
    1670                 1675                 1680

Gln Gly  Asn Gln Asp Ser Phe  Thr Pro Val Val Asn  Ser Leu Asp
    1685                 1690                 1695

Pro Pro  Leu Leu Thr Arg Tyr  Leu Arg Ile His Pro  Gln Ser Trp
    1700                 1705                 1710

Val His  Gln Ile Ala Leu Arg  Met Glu Val Leu Gly  Cys Glu Ala
    1715                 1720                 1725

Gln Asp  Leu Tyr Gly Ala Pro  Pro Gly Ala Ser Pro  Gly Thr Ser
    1730                 1735                 1740

Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1745                 1750                 1755

Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly
    1760                 1765                 1770

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Thr Pro Gly
    1775                 1780                 1785

Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser Ser Thr  Pro Ser Gly
    1790                 1795                 1800

Ala Thr  Gly Ser Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
    1805                 1810                 1815

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly
    1820                 1825                 1830

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Pro
    1835                 1840                 1845

Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ser Ser Pro  Ser Ala Ser
    1850                 1855                 1860

Thr Gly  Thr Gly Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1865                 1870                 1875

Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly
    1880                 1885                 1890

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Pro
    1895                 1900                 1905

Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ser Ser Pro  Ser Ala Ser
    1910                 1915                 1920
```

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
1925 1930 1935

Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
1940 1945 1950

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
1955 1960 1965

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser
1970 1975 1980

Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr
1985 1990 1995

Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
2000 2005 2010

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Ala Ser Ser Glu Gln
2015 2020 2025

Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala Thr Gly His His
2030 2035 2040

His His His His
2045

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<211> 6144
<212> DNA
<213> Artificial Sequence

<220>
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atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

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ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

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<210> 841
<211> 1903
<212> PRT
<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic polypeptide

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1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300
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Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575
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Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Thr Pro Gly
            740                 745                 750

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
        755                 760                 765

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    770                 775                 780

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
785                 790                 795                 800

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
                805                 810                 815

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            820                 825                 830

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
        835                 840                 845
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Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
    850                 855                 860

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
865                 870                 875                 880

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110
```

```
Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365
```

```
Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                1435                1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                1450                1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                1465                1470

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                1480                1485

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1490                1495                1500

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                1510                1515

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1520                1525                1530

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1535                1540                1545

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1550                1555                1560

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1565                1570                1575

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Pro Gly
    1580                1585                1590

Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser Pro Gly  Ala Ser Pro
    1595                1600                1605

Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Thr Pro Gly  Ser Gly Thr
    1610                1615                1620
```

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Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly
    1625                1630                1635

Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly
    1640                1645                1650

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Thr Pro Gly
    1655                1660                1665

Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser Ser Thr  Pro Ser Gly
    1670                1675                1680

Ala Thr  Gly Ser Pro Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly
    1685                1690                1695

Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr Gly Thr  Gly Pro Gly
    1700                1705                1710

Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly Pro Gly  Ala Ser Pro
    1715                1720                1725

Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Thr Pro Gly  Ser Gly Thr
    1730                1735                1740

Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly
    1745                1750                1755

Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr Gly Thr  Gly Pro Gly
    1760                1765                1770

Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly Pro Gly  Ala Ser Pro
    1775                1780                1785

Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1790                1795                1800

Ser Thr  Gly Ser Pro Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly
    1805                1810                1815

Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr Gly Thr  Gly Pro Gly
    1820                1825                1830

Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser Pro Gly  Ser Ser Pro
    1835                1840                1845

Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Thr Pro Gly  Ser Gly Thr
    1850                1855                1860

Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly
    1865                1870                1875
```

Ser Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
1880 1885 1890

Pro Ala Thr Gly His His His His His His
1895 1900

<210> 842
<211> 5712
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

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ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

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gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

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gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

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acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

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ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

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<210> 843
<211> 2047
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 843
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

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His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480
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Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Pro Gly Ala Ser
            740                 745                 750
```

```
Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
        755                 760                 765

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
    770                 775                 780

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro
785                 790                 795                 800

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
                805                 810                 815

Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
            820                 825                 830

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
        835                 840                 845

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
    850                 855                 860

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
865                 870                 875                 880

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
                885                 890                 895

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
            900                 905                 910

Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
        915                 920                 925

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
    930                 935                 940

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
945                 950                 955                 960

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
                965                 970                 975

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
            980                 985                 990

Pro Gly Thr Ser Ser Thr Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser
        995                 1000                1005

Thr Gly  Thr Gly Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
    1010                1015                1020
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Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Ala Ser
    1025                1030                1035

Ser Pro  Pro Val Leu Lys Arg  His Gln Ala Glu Ile  Thr Arg Thr
    1040                1045                1050

Thr Leu  Gln Ser Asp Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile
    1055                1060                1065

Ser Val  Glu Met Lys Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp
    1070                1075                1080

Glu Asn  Gln Ser Pro Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr
    1085                1090                1095

Phe Ile  Ala Ala Val Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser
    1100                1105                1110

Ser Pro  His Val Leu Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro
    1115                1120                1125

Gln Phe  Lys Lys Val Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe
    1130                1135                1140

Thr Gln  Pro Leu Tyr Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu
    1145                1150                1155

Leu Gly  Pro Tyr Ile Arg Ala  Glu Val Glu Asp Asn  Ile Met Val
    1160                1165                1170

Thr Phe  Arg Asn Gln Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser
    1175                1180                1185

Leu Ile  Ser Tyr Glu Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg
    1190                1195                1200

Lys Asn  Phe Val Lys Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys
    1205                1210                1215

Val Gln  His His Met Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys
    1220                1225                1230

Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu Glu Lys  Asp Val His
    1235                1240                1245

Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys His Thr  Asn Thr Leu
    1250                1255                1260

Asn Pro  Ala His Gly Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu
    1265                1270                1275
```

```
Phe Phe  Thr Ile Phe Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu
    1280                1285                1290

Asn Met  Glu Arg Asn Cys Arg  Ala Pro Cys Asn Ile  Gln Met Glu
    1295                1300                1305

Asp Pro  Thr Phe Lys Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly
    1310                1315                1320

Tyr Ile  Met Asp Thr Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln
    1325                1330                1335

Arg Ile  Arg Trp Tyr Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile
    1340                1345                1350

His Ser  Ile His Phe Ser Gly  His Val Phe Thr Val  Arg Lys Lys
    1355                1360                1365

Glu Glu  Tyr Lys Met Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe
    1370                1375                1380

Glu Thr  Val Glu Met Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val
    1385                1390                1395

Glu Cys  Leu Ile Gly Glu His  Leu His Ala Gly Met  Ser Thr Leu
    1400                1405                1410

Phe Leu  Val Tyr Ser Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala
    1415                1420                1425

Ser Gly  His Ile Arg Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr
    1430                1435                1440

Gly Gln  Trp Ala Pro Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser
    1445                1450                1455

Ile Asn  Ala Trp Ser Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val
    1460                1465                1470

Asp Leu  Leu Ala Pro Met Ile  Ile His Gly Ile Lys  Thr Gln Gly
    1475                1480                1485

Ala Arg  Gln Lys Phe Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile
    1490                1495                1500

Met Tyr  Ser Leu Asp Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn
    1505                1510                1515

Ser Thr  Gly Thr Leu Met Val  Phe Phe Gly Asn Val  Asp Ser Ser
    1520                1525                1530
```

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Gly Ile  Lys His Asn Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr
    1535                1540                1545

Ile Arg  Leu His Pro Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg
    1550                1555                1560

Met Glu  Leu Met Gly Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu
    1565                1570                1575

Gly Met  Glu Ser Lys Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser
    1580                1585                1590

Ser Tyr  Phe Thr Asn Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala
    1595                1600                1605

Arg Leu  His Leu Gln Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val
    1610                1615                1620

Asn Asn  Pro Lys Glu Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met
    1625                1630                1635

Lys Val  Thr Gly Val Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr
    1640                1645                1650

Ser Met  Tyr Val Lys Glu Phe  Leu Ile Ser Ser Ser  Gln Asp Gly
    1655                1660                1665

His Gln  Trp Thr Leu Phe Phe  Gln Asn Gly Lys Val  Lys Val Phe
    1670                1675                1680

Gln Gly  Asn Gln Asp Ser Phe  Thr Pro Val Val Asn  Ser Leu Asp
    1685                1690                1695

Pro Pro  Leu Leu Thr Arg Tyr  Leu Arg Ile His Pro  Gln Ser Trp
    1700                1705                1710

Val His  Gln Ile Ala Leu Arg  Met Glu Val Leu Gly  Cys Glu Ala
    1715                1720                1725

Gln Asp  Leu Tyr Gly Ala Pro  Pro Gly Ala Ser Pro  Gly Thr Ser
    1730                1735                1740

Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1745                1750                1755

Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly
    1760                1765                1770

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Thr Pro Gly
    1775                1780                1785
```

```
Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser Ser Thr  Pro Ser Gly
    1790                 1795                 1800

Ala Thr  Gly Ser Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
    1805                 1810                 1815

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly
    1820                 1825                 1830

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Pro
    1835                 1840                 1845

Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ser Ser Pro  Ser Ala Ser
    1850                 1855                 1860

Thr Gly  Thr Gly Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1865                 1870                 1875

Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly
    1880                 1885                 1890

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Pro
    1895                 1900                 1905

Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ser Ser Pro  Ser Ala Ser
    1910                 1915                 1920

Thr Gly  Thr Gly Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1925                 1930                 1935

Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly
    1940                 1945                 1950

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Pro
    1955                 1960                 1965

Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ala Ser Pro  Gly Thr Ser
    1970                 1975                 1980

Ser Thr  Gly Ser Pro Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr
    1985                 1990                 1995

Gly Pro  Gly Thr Pro Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly
    2000                 2005                 2010

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Ala Ser  Ser Glu Gln
    2015                 2020                 2025

Lys Leu  Ile Ser Glu Glu Asp  Leu Ser Pro Ala Thr  Gly His His
    2030                 2035                 2040
```

His His His His
    2045

<210> 844
<211> 6144
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
      polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

agagatctag cttcaggact cattggccct ctcctcatct gctacaaaga atctgtagat 1680

caaagaggaa accagataat gtcagacaag aggaatgtca tcctgttttc tgtatttgat 1740

gagaaccgaa gctggtacct cacagagaat atacaacgct ttctccccaa tccagctgga 1800

gtgcagcttg aggatccaga gttccaagcc tccaacatca tgcacagcat caatggctat 1860

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ccatatataa gagcagaagt tgaagataat atcatggtaa ctttcagaaa tcaggcctct 3540

cgtccctatt ccttctattc tagccttatt tcttatgagg aagatcagag gcaaggagca 3600

gaacctagaa aaaactttgt caagcctaat gaaaccaaaa cttacttttg gaaagtgcaa 3660

catcatatgg cacccactaa agatgagttt gactgcaaag cctgggctta tttctctgat 3720

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accgggtcag cctcgagcga acaaaaactc atctcagaag aggatctgtc acctgcaacc 6120

ggtcatcatc accatcacca ttga 6144

<210> 845
<211> 2047
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 845

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

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His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430
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Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700
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Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Thr Ser Glu
            740                 745                 750

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
        755                 760                 765

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
    770                 775                 780

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
785                 790                 795                 800

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
                805                 810                 815

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
            820                 825                 830

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
        835                 840                 845

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
    850                 855                 860

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
865                 870                 875                 880

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
                885                 890                 895

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
            900                 905                 910

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        915                 920                 925

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
    930                 935                 940

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
945                 950                 955                 960

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
                965                 970                 975
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Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
            980                 985                 990

Glu Pro Ser Glu Gly Ser Ala Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
        995                 1000                1005

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1010                1015                1020

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Ala Ser
    1025                1030                1035

Ser Pro  Pro Val Leu Lys Arg  His Gln Ala Glu Ile  Thr Arg Thr
    1040                1045                1050

Thr Leu  Gln Ser Asp Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile
    1055                1060                1065

Ser Val  Glu Met Lys Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp
    1070                1075                1080

Glu Asn  Gln Ser Pro Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr
    1085                1090                1095

Phe Ile  Ala Ala Val Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser
    1100                1105                1110

Ser Pro  His Val Leu Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro
    1115                1120                1125

Gln Phe  Lys Lys Val Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe
    1130                1135                1140

Thr Gln  Pro Leu Tyr Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu
    1145                1150                1155

Leu Gly  Pro Tyr Ile Arg Ala  Glu Val Glu Asp Asn  Ile Met Val
    1160                1165                1170

Thr Phe  Arg Asn Gln Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser
    1175                1180                1185

Leu Ile  Ser Tyr Glu Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg
    1190                1195                1200

Lys Asn  Phe Val Lys Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys
    1205                1210                1215

Val Gln  His His Met Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys
    1220                1225                1230
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Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu Glu Lys  Asp Val His
    1235                1240                1245

Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys His Thr  Asn Thr Leu
    1250                1255                1260

Asn Pro  Ala His Gly Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu
    1265                1270                1275

Phe Phe  Thr Ile Phe Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu
    1280                1285                1290

Asn Met  Glu Arg Asn Cys Arg  Ala Pro Cys Asn Ile  Gln Met Glu
    1295                1300                1305

Asp Pro  Thr Phe Lys Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly
    1310                1315                1320

Tyr Ile  Met Asp Thr Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln
    1325                1330                1335

Arg Ile  Arg Trp Tyr Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile
    1340                1345                1350

His Ser  Ile His Phe Ser Gly  His Val Phe Thr Val  Arg Lys Lys
    1355                1360                1365

Glu Glu  Tyr Lys Met Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe
    1370                1375                1380

Glu Thr  Val Glu Met Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val
    1385                1390                1395

Glu Cys  Leu Ile Gly Glu His  Leu His Ala Gly Met  Ser Thr Leu
    1400                1405                1410

Phe Leu  Val Tyr Ser Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala
    1415                1420                1425

Ser Gly  His Ile Arg Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr
    1430                1435                1440

Gly Gln  Trp Ala Pro Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser
    1445                1450                1455

Ile Asn  Ala Trp Ser Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val
    1460                1465                1470

Asp Leu  Leu Ala Pro Met Ile  Ile His Gly Ile Lys  Thr Gln Gly
    1475                1480                1485
```

```
Ala Arg  Gln Lys Phe Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile
    1490                1495                1500

Met Tyr  Ser Leu Asp Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn
    1505                1510                1515

Ser Thr  Gly Thr Leu Met Val  Phe Phe Gly Asn Val  Asp Ser Ser
    1520                1525                1530

Gly Ile  Lys His Asn Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr
    1535                1540                1545

Ile Arg  Leu His Pro Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg
    1550                1555                1560

Met Glu  Leu Met Gly Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu
    1565                1570                1575

Gly Met  Glu Ser Lys Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser
    1580                1585                1590

Ser Tyr  Phe Thr Asn Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala
    1595                1600                1605

Arg Leu  His Leu Gln Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val
    1610                1615                1620

Asn Asn  Pro Lys Glu Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met
    1625                1630                1635

Lys Val  Thr Gly Val Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr
    1640                1645                1650

Ser Met  Tyr Val Lys Glu Phe  Leu Ile Ser Ser Ser  Gln Asp Gly
    1655                1660                1665

His Gln  Trp Thr Leu Phe Phe  Gln Asn Gly Lys Val  Lys Val Phe
    1670                1675                1680

Gln Gly  Asn Gln Asp Ser Phe  Thr Pro Val Val Asn  Ser Leu Asp
    1685                1690                1695

Pro Pro  Leu Leu Thr Arg Tyr  Leu Arg Ile His Pro  Gln Ser Trp
    1700                1705                1710

Val His  Gln Ile Ala Leu Arg  Met Glu Val Leu Gly  Cys Glu Ala
    1715                1720                1725

Gln Asp  Leu Tyr Gly Ala Pro  Pro Gly Ala Ser Pro  Gly Thr Ser
    1730                1735                1740
```

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Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1745                1750                1755

Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly
    1760                1765                1770

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Thr Pro Gly
    1775                1780                1785

Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser Ser Thr  Pro Ser Gly
    1790                1795                1800

Ala Thr  Gly Ser Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
    1805                1810                1815

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly
    1820                1825                1830

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Pro
    1835                1840                1845

Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ser Ser Pro  Ser Ala Ser
    1850                1855                1860

Thr Gly  Thr Gly Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1865                1870                1875

Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly
    1880                1885                1890

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Pro
    1895                1900                1905

Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ser Ser Pro  Ser Ala Ser
    1910                1915                1920

Thr Gly  Thr Gly Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1925                1930                1935

Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly
    1940                1945                1950

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Pro
    1955                1960                1965

Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ala Ser Pro  Gly Thr Ser
    1970                1975                1980

Ser Thr  Gly Ser Pro Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr
    1985                1990                1995
```

Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
2000 2005 2010

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Ala Ser Ser Glu Gln
2015 2020 2025

Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala Thr Gly His His
2030 2035 2040

His His His His
2045

<210> 846
<211> 6144
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

agagatctag cttcaggact cattggccct ctcctcatct gctacaaaga atctgtagat 1680

caaagaggaa accagataat gtcagacaag aggaatgtca tcctgttttc tgtatttgat 1740

gagaaccgaa gctggtacct cacagagaat atacaacgct ttctccccaa tccagctgga 1800

gtgcagcttg aggatccaga gttccaagcc tccaacatca tgcacagcat caatggctat 1860

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agggctcaga gtggcagtgt ccctcagttc aagaaagttg ttttccagga atttactgat 3420

ggctccttta ctcagccctt ataccgtgga gaactaaatg aacatttggg actcctgggg 3480

ccatatataa gagcagaagt tgaagataat atcatggtaa ctttcagaaa tcaggcctct 3540

cgtccctatt ccttctattc tagccttatt tcttatgagg aagatcagag gcaaggagca 3600

gaacctagaa aaaactttgt caagcctaat gaaaccaaaa cttacttttg gaaagtgcaa 3660

catcatatgg cacccactaa agatgagttt gactgcaaag cctgggctta tttctctgat 3720

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gctccctgca atatccagat ggaagatccc acttttaaag agaattatcg cttccatgca 3960

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<210> 847
<211> 2047
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 847
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

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Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380
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Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655
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Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Pro Gly Ala Ser
            740                 745                 750

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
        755                 760                 765

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
    770                 775                 780

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro
785                 790                 795                 800

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
                805                 810                 815

Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
            820                 825                 830

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
        835                 840                 845

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
    850                 855                 860

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
865                 870                 875                 880

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
                885                 890                 895

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
            900                 905                 910

Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
        915                 920                 925
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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
    930                 935                 940

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
945                 950                 955                 960

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
                965                 970                 975

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
            980                 985                 990

Pro Gly Thr Ser Ser Thr Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser
        995                 1000                1005

Thr Gly  Thr Gly Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
    1010                1015                1020

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Ala Ser
    1025                1030                1035

Ser Pro  Pro Val Leu Lys Arg  His Gln Arg Glu Ile  Thr Arg Thr
    1040                1045                1050

Thr Leu  Gln Ser Asp Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile
    1055                1060                1065

Ser Val  Glu Met Lys Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp
    1070                1075                1080

Glu Asn  Gln Ser Pro Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr
    1085                1090                1095

Phe Ile  Ala Ala Val Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser
    1100                1105                1110

Ser Pro  His Val Leu Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro
    1115                1120                1125

Gln Phe  Lys Lys Val Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe
    1130                1135                1140

Thr Gln  Pro Leu Tyr Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu
    1145                1150                1155

Leu Gly  Pro Tyr Ile Arg Ala  Glu Val Glu Asp Asn  Ile Met Val
    1160                1165                1170

Thr Phe  Arg Asn Gln Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser
    1175                1180                1185
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Leu Ile  Ser Tyr Glu Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg
    1190                 1195                 1200

Lys Asn  Phe Val Lys Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys
    1205                 1210                 1215

Val Gln  His His Met Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys
    1220                 1225                 1230

Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu Glu Lys  Asp Val His
    1235                 1240                 1245

Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys His Thr  Asn Thr Leu
    1250                 1255                 1260

Asn Pro  Ala His Gly Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu
    1265                 1270                 1275

Phe Phe  Thr Ile Phe Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu
    1280                 1285                 1290

Asn Met  Glu Arg Asn Cys Arg  Ala Pro Cys Asn Ile  Gln Met Glu
    1295                 1300                 1305

Asp Pro  Thr Phe Lys Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly
    1310                 1315                 1320

Tyr Ile  Met Asp Thr Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln
    1325                 1330                 1335

Arg Ile  Arg Trp Tyr Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile
    1340                 1345                 1350

His Ser  Ile His Phe Ser Gly  His Val Phe Thr Val  Arg Lys Lys
    1355                 1360                 1365

Glu Glu  Tyr Lys Met Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe
    1370                 1375                 1380

Glu Thr  Val Glu Met Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val
    1385                 1390                 1395

Glu Cys  Leu Ile Gly Glu His  Leu His Ala Gly Met  Ser Thr Leu
    1400                 1405                 1410

Phe Leu  Val Tyr Ser Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala
    1415                 1420                 1425

Ser Gly  His Ile Arg Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr
    1430                 1435                 1440
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Gly Gln  Trp Ala Pro Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser
    1445                1450                1455

Ile Asn  Ala Trp Ser Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val
    1460                1465                1470

Asp Leu  Leu Ala Pro Met Ile  Ile His Gly Ile Lys  Thr Gln Gly
    1475                1480                1485

Ala Arg  Gln Lys Phe Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile
    1490                1495                1500

Met Tyr  Ser Leu Asp Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn
    1505                1510                1515

Ser Thr  Gly Thr Leu Met Val  Phe Phe Gly Asn Val  Asp Ser Ser
    1520                1525                1530

Gly Ile  Lys His Asn Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr
    1535                1540                1545

Ile Arg  Leu His Pro Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg
    1550                1555                1560

Met Glu  Leu Met Gly Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu
    1565                1570                1575

Gly Met  Glu Ser Lys Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser
    1580                1585                1590

Ser Tyr  Phe Thr Asn Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala
    1595                1600                1605

Arg Leu  His Leu Gln Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val
    1610                1615                1620

Asn Asn  Pro Lys Glu Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met
    1625                1630                1635

Lys Val  Thr Gly Val Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr
    1640                1645                1650

Ser Met  Tyr Val Lys Glu Phe  Leu Ile Ser Ser Ser  Gln Asp Gly
    1655                1660                1665

His Gln  Trp Thr Leu Phe Phe  Gln Asn Gly Lys Val  Lys Val Phe
    1670                1675                1680

Gln Gly  Asn Gln Asp Ser Phe  Thr Pro Val Val Asn  Ser Leu Asp
    1685                1690                1695
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Pro Pro  Leu Leu Thr Arg Tyr  Leu Arg Ile His Pro  Gln Ser Trp
    1700                1705                1710

Val His  Gln Ile Ala Leu Arg  Met Glu Val Leu Gly  Cys Glu Ala
    1715                1720                1725

Gln Asp  Leu Tyr Gly Ala Pro  Pro Gly Ala Ser Pro  Gly Thr Ser
    1730                1735                1740

Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1745                1750                1755

Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly
    1760                1765                1770

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Thr Pro Gly
    1775                1780                1785

Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser Ser Thr  Pro Ser Gly
    1790                1795                1800

Ala Thr  Gly Ser Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
    1805                1810                1815

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly
    1820                1825                1830

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Pro
    1835                1840                1845

Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ser Ser Pro  Ser Ala Ser
    1850                1855                1860

Thr Gly  Thr Gly Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1865                1870                1875

Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly
    1880                1885                1890

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Pro
    1895                1900                1905

Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ser Ser Pro  Ser Ala Ser
    1910                1915                1920

Thr Gly  Thr Gly Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1925                1930                1935

Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly
    1940                1945                1950
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Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
1955 1960 1965

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser
1970 1975 1980

Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr
1985 1990 1995

Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
2000 2005 2010

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Ala Ser Ser Glu Gln
2015 2020 2025

Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala Thr Gly His His
2030 2035 2040

His His His His
2045

<210> 848
<211> 6144
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

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catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

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tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

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catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

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gagaaccgaa gctggtacct cacagagaat atacaacgct ttctccccaa tccagctgga 1800

gtgcagcttg aggatccaga gttccaagcc tccaacatca tgcacagcat caatggctat 1860

gtttttgata gtttgcagtt gtcagtttgt ttgcatgagg tggcatactg gtacattcta 1920

agcattggag cacagactga cttcctttct gtcttcttct ctggatatac cttcaaacac 1980

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ggtcatcatc accatcacca ttga 6144

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<211> 1903
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val

65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp

340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser

610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Thr Pro Gly
740 745 750

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
755 760 765

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
770 775 780

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
785 790 795 800

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
805 810 815

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
820 825 830

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
835 840 845

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
850 855 860

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
865 870 875 880

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Pro

885 890 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
900 905 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
915 920 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
930 935 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945 950 955 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
965 970 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
980 985 990

Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
995 1000 1005

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1010 1015 1020

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1025 1030 1035

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1040 1045 1050

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1055 1060 1065

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1070 1075 1080

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1085 1090 1095

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1100 1105 1110

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1115 1120 1125

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1130 1135 1140

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro

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Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
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Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1415 1420 1425

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1430 1435 1440

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1445 1450 1455

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1460 1465 1470

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1475 1480 1485

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1490 1495 1500

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1505 1510 1515

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1520 1525 1530

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Pro Gly Thr
1580 1585 1590

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1595 1600 1605

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
1610 1615 1620

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1625 1630 1635

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1640 1645 1650

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly

1655 1660 1665

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
1670 1675 1680

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1685 1690 1695

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
1700 1705 1710

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
1715 1720 1725

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
1730 1735 1740

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1745 1750 1755

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1760 1765 1770

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1775 1780 1785

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
1790 1795 1800

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
1805 1810 1815

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
1820 1825 1830

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
1835 1840 1845

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
1850 1855 1860

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
1865 1870 1875

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
1880 1885 1890

Pro Ala Thr Gly His His His His His His
1895 1900

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<211> 5712
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

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ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

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gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

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<210> 851
<211> 1769

<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 851

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Gly Ser Ser
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Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
        35                  40                  45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
    50                  55                  60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65                  70                  75                  80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
                85                  90                  95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
            100                 105                 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
        115                 120                 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
    130                 135                 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145                 150                 155                 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240
```

```
Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510
```

```
Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu
545                 550                 555                 560

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
                565                 570                 575

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            580                 585                 590

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
        595                 600                 605

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    610                 615                 620

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
625                 630                 635                 640

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
                645                 650                 655

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
            660                 665                 670

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
        675                 680                 685

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp
    690                 695                 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780
```

```
Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020

Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050
```

```
Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305
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Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
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Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                 1330                 1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                 1345                 1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                 1360                 1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                 1375                 1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                 1390                 1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                 1405                 1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                 1420                 1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                 1435                 1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                 1450                 1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                 1465                 1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                 1480                 1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                 1495                 1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                 1510                 1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                 1525                 1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                 1540                 1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                 1555                 1560
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Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
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Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
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Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
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Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1610 1615 1620

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
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Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
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Ser Pro Ala Thr Gly His His His His His His
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<211> 5310
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<223> Description of Artificial Sequence: Synthetic polynucleotide

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<212> PRT
<213> Artificial Sequence

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Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
35 40 45

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
50 55 60

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu

65 70 75 80

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
85 90 95

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
100 105 110

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
115 120 125

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
130 135 140

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
145 150 155 160

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ala Ser Ser
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
260 265 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
275 280 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
290 295 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305 310 315 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
325 330 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe

340 345 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
355 360 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
370 375 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385 390 395 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
405 410 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
420 425 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
435 440 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
450 455 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465 470 475 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
485 490 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
500 505 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
515 520 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
530 535 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu
545 550 555 560

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
565 570 575

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
580 585 590

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
595 600 605

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly

610 615 620

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
625 630 635 640

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
645 650 655

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
660 665 670

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
675 680 685

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp
690 695 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705 710 715 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
725 730 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
740 745 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
755 760 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
770 775 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785 790 795 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
805 810 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
820 825 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
835 840 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
850 855 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865 870 875 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro

885 890 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
900 905 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
915 920 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
930 935 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945 950 955 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
965 970 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
980 985 990

His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val
995 1000 1005

Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr
1010 1015 1020

Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu
1025 1030 1035

Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His Gln
1040 1045 1050

Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
1055 1060 1065

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
1070 1075 1080

Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln
1085 1090 1095

Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp
1100 1105 1110

Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala
1115 1120 1125

Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu
1130 1135 1140

Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu

1145 1150 1155

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1160 1165 1170

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1175 1180 1185

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1190 1195 1200

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1205 1210 1215

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1220 1225 1230

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1235 1240 1245

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1250 1255 1260

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1265 1270 1275

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1280 1285 1290

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1295 1300 1305

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1310 1315 1320

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1325 1330 1335

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1340 1345 1350

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1355 1360 1365

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1370 1375 1380

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1385 1390 1395

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His

1400 1405 1410

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1415 1420 1425

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1430 1435 1440

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1445 1450 1455

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1460 1465 1470

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1475 1480 1485

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1490 1495 1500

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1505 1510 1515

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1520 1525 1530

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1535 1540 1545

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1550 1555 1560

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1565 1570 1575

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1580 1585 1590

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1595 1600 1605

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1610 1615 1620

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1625 1630 1635

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile

1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
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Ser Pro Ala Thr Gly His His His His His His
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aactctctag acccaccgtt actgactcgc taccttcgaa ttcaccccca gagttgggtg 5160

caccagattg ccctgaggat ggaggttctg ggctgcgagg cacaggacct ctacggcgcc 5220

ggatcacctg gggccgaaac ggccgaacaa aaactcatct cagaagagga tctgtcacct 5280

gcaaccggtc atcatcacca tcaccattga 5310

<210> 855
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 855
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Gly Ala Pro Gly Ser Ser Pro Ser
35 40 45

Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
50 55 60

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65 70 75 80

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro
85 90 95

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
100 105 110

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
115 120 125

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser
130 135 140

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145 150 155 160

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
165 170 175

```
Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445
```

```
Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu
545                 550                 555                 560

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
                565                 570                 575

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            580                 585                 590

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
        595                 600                 605

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    610                 615                 620

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
625                 630                 635                 640

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
                645                 650                 655

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
            660                 665                 670

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
        675                 680                 685

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp
    690                 695                 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720
```

```
Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990
```

```
His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020

Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245
```

```
Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500
```

```
Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1730                1735                1740

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
    1745                1750                1755
```

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 856
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 856
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ggcgcgccag gctccagccc ctccgcgagc acgggaaccg gaccaggttc gtcaccctca 180

gcatcaacgg ggacgggacc gggggcgtca ccaggaacgt cctccaccgg ctcgccgggt 240

gcatcacccg gaacgtcatc gaccggatcg ccagggagct cgacgccatc aggcgcaaca 300

ggatcacctg gctcaagccc tagcgcgtca accggcacgg gtccgggtgc ctcccctggc 360

acgtccagca ccggatcacc cggatcgagc ccatccgcct caaccggaac cggacccggt 420

acaccagggt cgggaacagc ctcctcgtca ccaggctcct caacccctc gggagccacg 480

ggttcgcccg gttcgtcaac gccttccgga gcaactggta gccccggagc atcgccagga 540

acttcgagca cggggtcgcc cgcctcgagc aacacctcag tcgtgtacaa aaagactctg 600

tttgtagaat tcacggatca ccttttcaac atcgctaagc caaggccacc ctggatgggt 660

ctgctaggtc ctaccatcca ggctgaggtt tatgatacag tggtcattac acttaagaac 720

atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag 780

ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct 840

ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

ggcctcattg gagccctact agtatgtaga gaagggagtc tggccaagga aaagacacag 1020

accttgcaca aatttatact acttttgct gtatttgatg aagggaaaag ttggcactca 1080

gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg 1200

aaatcagtct attggcatgt gattggaatg ggcaccactc ctgaagtgca ctcaatattc 1260

ctcgaaggtc acacatttct tgtgaggaac catcgccagg ctagcttgga aatctcgcca 1320

ataactttcc ttactgctca aacactcttg atggaccttg gacagtttct actgttttgt 1380

catatctctt cccaccaaca tgatggcatg gaagcttatg tcaaagtaga cagctgtcca 1440

gaggaacccc aactacgaat gaaaaataat gaagaagcgg aagactatga tgatgatctt 1500

actgattctg aaatggatgt ggtcaggttt gatgatgaca actctccttc ctttatccaa 1560

attcgctcag ttgccaagaa gcatcctaaa acttgggtac attacattgc tgctgaagag 1620

gaggactggg actatgctcc cttagtcctc gcccccgatg gcgcgccaac cagtacggag 1680

ccgtccgagg ggagcgcacc aggaagcccg gctgggagcc cgacttctac cgaagagggt 1740

acatctaccg aaccaagtga aggttcagca ccaggcacct caacagaacc ctctgagggc 1800

tcggcgcctg gtacaagtga gtccgccacc ccagaatccg ggcctgggac aagcacagaa 1860

ccttcggaag ggagtgcccc tggaacatcc gaatcggcaa ccccagaatc agggccagga 1920

tctgagcccg cgacttcggg ctccgagacg cctgggacat ccaccgagcc ctccgaagga 1980

tcagccccag gcaccagcac ggagccctct gagggaagcg cacctggtac cagcgaaagc 2040

gcaactcccg aatcaggtcc cggtacgagc gagtcggcga ccccggagag cgggccaggt 2100

gcctcgagcg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 2160

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 2220

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 2280

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 2340

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 2400

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 2460

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 2520

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ggcatgaccg ccttactgaa ggtttctagt tgtgacaaga acactggtga ttattacgag 3060

gacagttatg aagatatttc agcatacttg ctgagtaaaa acaatgccat tgaaccaaga 3120

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gatcagaggc aaggagcaga acctagaaaa aactttgtca agcctaatga aaccaaaact 3660

tacttttgga aagtgcaaca tcatatggca cccactaaag atgagtttga ctgcaaagcc 3720

tgggcttatt tctctgatgt tgacctggaa aaagatgtgc actcaggcct gattggaccc 3780

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gaatttgctc tgtttttcac catctttgat gagaccaaaa gctggtactt cactgaaaat 3900

atggaaagaa actgcagggc tccctgcaat atccagatgg aagatcccac ttttaaagag 3960

aattatcgct tccatgcaat caatggctac ataatggata cactacctgg cttagtaatg 4020

gctcaggatc aaaggattcg atggtatctg ctcagcatgg gcagcaatga aaacatccat 4080

tctattcatt tcagtggaca tgtgttcact gtacgaaaaa aagaggagta taaaatggca 4140

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atttggcggg tggaatgcct tattggcgag catctacatg ctgggatgag cacacttttt 4260

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cattatagca ttcgcagcac tcttcgcatg gagttgatgg gctgtgattt aaatagttgc 4740

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atgtatgtga aggagttcct catctccagc agtcaagatg gccatcagtg gactctcttt 5040

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caccagattg ccctgaggat ggaggttctg ggctgcgagg cacaggacct ctacggcgcc 5220

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gcaaccggtc atcatcacca tcaccattga 5310

<210> 857
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 857

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Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1               5                   10                  15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Gly Ala Pro Thr Ser Glu Ser Ala
        35                  40                  45

Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu
    50                  55                  60

Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
65                  70                  75                  80

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala
                85                  90                  95

Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
            100                 105                 110

Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr
        115                 120                 125

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr
    130                 135                 140

Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
145                 150                 155                 160

Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser
                165                 170                 175

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ala Ser Ser Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270
```

```
Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540
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Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu
545                 550                 555                 560

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
                565                 570                 575

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            580                 585                 590

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
        595                 600                 605

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    610                 615                 620

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
625                 630                 635                 640

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
                645                 650                 655

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
            660                 665                 670

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
        675                 680                 685

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp
    690                 695                 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815
```

```
Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020

Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080
```

```
Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                 1090                 1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                 1105                 1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                 1120                 1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                 1135                 1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                 1150                 1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                 1165                 1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                 1180                 1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                 1195                 1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                 1210                 1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                 1225                 1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                 1240                 1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                 1255                 1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                 1270                 1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                 1285                 1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                 1300                 1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                 1315                 1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                 1330                 1335
```

```
Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590
```

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1595 1600 1605

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1610 1615 1620

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1625 1630 1635

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 858
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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ggcgcgccaa catcagagag cgccacccct gaaagtggtc ccgggagcga gccagccaca 180

tctgggtcgg aaacgccagg cacaagtgag tctgcaactc ccgagtccgg acctggctcc 240

gagcctgcca ctagcggctc cgagactccg ggaacttccg agagcgctac accagaaagc 300

ggacccggaa ccagtaccga acctagcgag ggctctgctc cgggcagccc agccggctct 360
cctacatcca cggaggaggg cacttccgaa tccgccaccc cggagtcagg gccaggatct 420
gaacccgcta cctcaggcag tgagacgcca ggaacgagcg agtccgctac accggagagt 480
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tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 2460

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gctcaggatc aaaggattcg atggtatctg ctcagcatgg gcagcaatga aaacatccat 4080

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<210> 859
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 859
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Gly Ala Pro Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
20 25 30

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
35 40 45

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
50 55 60

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
65 70 75 80

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
85 90 95

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Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
            100                 105                 110

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
        115                 120                 125

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
    130                 135                 140

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
145                 150                 155                 160

Ser Thr Glu Glu Gly Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365
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```
Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu
545                 550                 555                 560

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
                565                 570                 575

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            580                 585                 590

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
        595                 600                 605

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    610                 615                 620

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
625                 630                 635                 640
```

```
Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
                645                 650                 655

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
            660                 665                 670

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
        675                 680                 685

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp
    690                 695                 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910
```

```
Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020

Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170
```

```
Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425
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Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680
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Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 860
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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tcggaaacgc caggcacaag tgagtctgca actcccgagt ccggacctgg ctccgagcct 180

gccactagcg gctccgagac tccgggaact tccgagagcg ctacaccaga aagcggaccc 240

ggaaccagta ccgaacctag cgagggctct gctccgggca gcccagccgg ctctcctaca 300

tccacggagg agggcacttc cgaatccgcc accccggagt cagggccagg atctgaaccc 360

gctacctcag gcagtgagac gccaggaacg agcgagtccg ctacaccgga gagtgggcca 420

gggagccctg ctggatctcc tacgtccact gaggaagggt caccagcggg ctcgcccacc 480

agcactgaag aaggtgcctc gagcagtgat ctcggtgagc tgcctgtgga cgcaagattt 540

cctcctagag tgccaaaatc ttttccattc aacacctcag tcgtgtacaa aaagactctg 600

tttgtagaat tcacggatca ccttttcaac atcgctaagc caaggccacc ctggatgggt 660

ctgctaggtc ctaccatcca ggctgaggtt tatgatacag tggtcattac acttaagaac 720

atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag 780

ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct 840

ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

ggcctcattg gagccctact agtatgtaga gaagggagtc tggccaagga aaagacacag 1020

accttgcaca aatttatact acttttgct gtatttgatg aagggaaaag ttggcactca 1080

gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg 1200

aaatcagtct attggcatgt gattggaatg ggcaccactc ctgaagtgca ctcaatattc 1260

ctcgaaggtc acacatttct tgtgaggaac catcgccagg ctagcttgga aatctcgcca 1320

ataactttcc ttactgctca aacactcttg atggaccttg gacagtttct actgttttgt 1380

catatctctt cccaccaaca tgatggcatg gaagcttatg tcaaagtaga cagctgtcca 1440

gaggaacccc aactacgaat gaaaaataat gaagaagcgg aagactatga tgatgatctt 1500

actgattctg aaatggatgt ggtcaggttt gatgatgaca actctccttc ctttatccaa 1560

attcgctcag ttgccaagaa gcatcctaaa acttgggtac attacattgc tgctgaagag 1620

gaggactggg actatgctcc cttagtcctc gcccccgatg gcgcgccaac cagtacggag 1680

ccgtccgagg ggagcgcacc aggaagcccg gctgggagcc cgacttctac cgaagagggt 1740

acatctaccg aaccaagtga aggttcagca ccaggcacct caacagaacc ctctgagggc 1800

tcggcgcctg gtacaagtga gtccgccacc ccagaatccg ggcctgggac aagcacagaa 1860

ccttcggaag ggagtgcccc tggaacatcc gaatcggcaa ccccagaatc agggccagga 1920

tctgagcccg cgacttcggg ctccgagacg cctgggacat ccaccgagcc ctccgaagga 1980

tcagccccag gcaccagcac ggagccctct gagggaagcg cacctggtac cagcgaaagc 2040

gcaactcccg aatcaggtcc cggtacgagc gagtcggcga ccccggagag cgggccaggt 2100

gcctcgagcg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 2160

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 2220

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 2280

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 2340

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 2400

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caaagaggaa accagataat gtcagacaag aggaatgtca tcctgttttc tgtatttgat 2640

gagaaccgaa gctggtacct cacagagaat atacaacgct ttctccccaa tccagctgga 2700

gtgcagcttg aggatccaga gttccaagcc tccaacatca tgcacagcat caatggctat 2760

gtttttgata gtttgcagtt gtcagtttgt ttgcatgagg tggcatactg gtacattcta 2820

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ggcatgaccg ccttactgaa ggtttctagt tgtgacaaga acactggtga ttattacgag 3060

gacagttatg aagatatttc agcatacttg ctgagtaaaa acaatgccat tgaaccaaga 3120

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cttcagtcag atcaagagga aatcgattat gatgatacca tatcagttga aatgaagaag 3240

gaagattttg acatttatga tgaggatgaa aatcagagcc cccgcagctt tcaaaagaaa 3300

acacgacact attttattgc tgcagtggag aggctctggg attatgggat gagtagctcc 3360

ccacatgttc taagaaacag ggctcagagt ggcagtgtcc ctcagttcaa gaaagttgtt 3420

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catttgggac tcctggggcc atatataaga gcagaagttg aagataatat catggtaact 3540

ttcagaaatc aggcctctcg tccctattcc ttctattcta gccttatttc ttatgaggaa 3600

gatcagaggc aaggagcaga acctagaaaa aactttgtca agcctaatga aaccaaaact 3660

tacttttgga aagtgcaaca tcatatggca cccactaaag atgagtttga ctgcaaagcc 3720

tgggcttatt tctctgatgt tgacctggaa aaagatgtgc actcaggcct gattggaccc 3780

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gctcaggatc aaaggattcg atggtatctg ctcagcatgg gcagcaatga aaacatccat 4080

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atttggcggg tggaatgcct tattggcgag catctacatg ctgggatgag cacacttttt 4260

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cattatagca ttcgcagcac tcttcgcatg gagttgatgg gctgtgattt aaatagttgc 4740

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tactttacca atatgtttgc cacctggtct ccttcaaaag ctcgacttca cctccaaggg 4860

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cagaagacaa tgaaagtcac aggagtaact actcagggag taaaatctct gcttaccagc 4980

atgtatgtga aggagttcct catctccagc agtcaagatg gccatcagtg gactctcttt 5040

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caccagattg ccctgaggat ggaggttctg ggctgcgagg cacaggacct ctacggcgcc 5220

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gcaaccggtc atcatcacca tcaccattga 5310

<210> 861
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 861

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Thr Ser Glu
20 25 30

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
35 40 45

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
50 55 60

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
65 70 75 80

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
85 90 95

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
100 105 110

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
115 120 125

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
130 135 140

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
145 150 155 160

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ala Ser Ser
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu

195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
260 265 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
275 280 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
290 295 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305 310 315 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
325 330 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
340 345 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
355 360 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
370 375 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385 390 395 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
405 410 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
420 425 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
435 440 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
450 455 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro

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465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Ala Pro
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740 745 750

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
755 760 765

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
770 775 780

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
785 790 795 800

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
805 810 815

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
820 825 830

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
835 840 845

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
850 855 860

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
865 870 875 880

Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
885 890 895

Ala Ser Ser Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
900 905 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
915 920 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
930 935 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945 950 955 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
965 970 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
980 985 990

His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val
995 1000 1005

Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr

1010 1015 1020

Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu
1025 1030 1035

Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His Gln
1040 1045 1050

Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
1055 1060 1065

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
1070 1075 1080

Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln
1085 1090 1095

Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp
1100 1105 1110

Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala
1115 1120 1125

Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu
1130 1135 1140

Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
1145 1150 1155

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1160 1165 1170

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1175 1180 1185

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1190 1195 1200

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1205 1210 1215

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1220 1225 1230

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1235 1240 1245

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1250 1255 1260

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr

1265 1270 1275

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1280 1285 1290

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1295 1300 1305

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1310 1315 1320

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1325 1330 1335

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1340 1345 1350

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1355 1360 1365

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1370 1375 1380

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1385 1390 1395

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1400 1405 1410

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1415 1420 1425

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1430 1435 1440

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1445 1450 1455

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1460 1465 1470

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1475 1480 1485

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1490 1495 1500

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1505 1510 1515

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe

1520 1525 1530

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1535 1540 1545

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1550 1555 1560

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1565 1570 1575

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1580 1585 1590

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1595 1600 1605

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1610 1615 1620

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1625 1630 1635

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 862

<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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ctcggtgagc tgcctgtggg cgcgccaaca tcagagagcg ccacccctga aagtggtccc 120

gggagcgagc cagccacatc tgggtcggaa acgccaggca caagtgagtc tgcaactccc 180

gagtccggac ctggctccga gcctgccact agcggctccg agactccggg aacttccgag 240

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
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        35                  40                  45

Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
    50                  55                  60

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65                  70                  75                  80

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro
                85                  90                  95

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
            100                 105                 110

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
        115                 120                 125

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser
    130                 135                 140

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145                 150                 155                 160

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
                165                 170                 175

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300
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Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
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Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575
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Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Ala Pro
            740                 745                 750

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
        755                 760                 765

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
    770                 775                 780

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
785                 790                 795                 800

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
                805                 810                 815

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
            820                 825                 830

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
        835                 840                 845
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Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
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Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
865                 870                 875                 880

Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
                885                 890                 895

Ala Ser Ser Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020

Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
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Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
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Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
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Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
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Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365
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Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
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Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
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Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
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Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
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Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
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Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
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Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
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Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
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<212> PRT
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<400> 865

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Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
        35                  40                  45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
    50                  55                  60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65                  70                  75                  80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
                85                  90                  95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
            100                 105                 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
        115                 120                 125
```

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Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
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Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145                 150                 155                 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400
```

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Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
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His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670
```

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Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Ala Pro
            740                 745                 750

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
        755                 760                 765

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
    770                 775                 780

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
785                 790                 795                 800

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
                805                 810                 815

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
            820                 825                 830

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
        835                 840                 845

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
    850                 855                 860

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
865                 870                 875                 880

Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
                885                 890                 895

Ala Ser Ser Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940
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```
Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
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Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020

Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200
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Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
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Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
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Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                 1240                 1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                 1255                 1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                 1270                 1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                 1285                 1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                 1300                 1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                 1315                 1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                 1330                 1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                 1345                 1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                 1360                 1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                 1375                 1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                 1390                 1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                 1405                 1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                 1420                 1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                 1435                 1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                 1450                 1455
```

```
Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
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Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
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Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
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Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710
```

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
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Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
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Ser Pro Ala Thr Gly His His His His His His
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agcctctaca tctctcagtt tatcatcatg tatagtcttg atgggaagaa gtggcagact 4560
tatcgaggaa attccactgg aaccttaatg gtcttctttg gcaatgtgga ttcatctggg 4620
ataaaacaca atatttttaa ccctccaatt attgctcgat acatccgttt gcacccaact 4680
cattatagca ttcgcagcac tcttcgcatg gagttgatgg gctgtgattt aaatagttgc 4740
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<210> 867
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 867
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Met Gln Gly Ala Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
20 25 30

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
35 40 45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
50 55 60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65 70 75 80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
85 90 95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
100 105 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
115 120 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
130 135 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145 150 155 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

```
Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495
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Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Ala Pro
            740                 745                 750

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
        755                 760                 765
```

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Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
    770                 775                 780

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
785                 790                 795                 800

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
                805                 810                 815

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
            820                 825                 830

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
        835                 840                 845

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
    850                 855                 860

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
865                 870                 875                 880

Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
                885                 890                 895

Ala Ser Ser Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020

Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035
```

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Pro Arg  Ser Phe Ser Gln Asn  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290
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Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                 1300                 1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                 1315                 1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                 1330                 1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                 1345                 1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                 1360                 1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                 1375                 1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                 1390                 1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                 1405                 1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                 1420                 1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                 1435                 1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                 1450                 1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                 1465                 1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                 1480                 1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                 1495                 1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                 1510                 1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                 1525                 1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                 1540                 1545
```

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1550 1555 1560

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1565 1570 1575

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1580 1585 1590

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1595 1600 1605

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1610 1615 1620

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1625 1630 1635

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 868
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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acggggacgg gaccgggggc gtcaccagga acgtcctcca ccggctcgcc gggtgcatca 180

cccggaacgt catcgaccgg atcgccaggg agctcgacgc catcaggcgc aacaggatca 240

cctggctcaa gccctagcgc gtcaaccggc acgggtccgg gtgcctcccc tggcacgtcc 300

agcaccggat cacccggatc gagcccatcc gcctcaaccg gaaccggacc cggtacacca 360

gggtcgggaa cagcctcctc gtcaccaggc tcctcaaccc cctcgggagc cacgggttcg 420

cccggttcgt caacgccttc cggagcaact ggtagccccg gagcatcgcc aggaacttcg 480

agcacggggt cgcccgcctc gagcagtgat ctcggtgagc tgcctgtgga cgcaagattt 540

cctcctagag tgccaaaatc ttttccattc aacacctcag tcgtgtacaa aaagactctg 600

tttgtagaat tcacggatca ccttttcaac atcgctaagc caaggccacc ctggatgggt 660

ctgctaggtc ctaccatcca ggctgaggtt tatgatacag tggtcattac acttaagaac 720

atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag 780

ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct 840

ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

ggcctcattg gagccctact agtatgtaga gaagggagtc tggccaagga aaagacacag 1020

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gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg 1200

aaatcagtct attggcatgt gattggaatg ggcaccactc ctgaagtgca ctcaatattc 1260

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<210> 869
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 869
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Gly Ala Pro Gly Ser Ser Pro Ser
35 40 45

Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly

50 55 60

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65 70 75 80

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro
85 90 95

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
100 105 110

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
115 120 125

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser
130 135 140

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145 150 155 160

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
165 170 175

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
260 265 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
275 280 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
290 295 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305 310 315 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys

325 330 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
340 345 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
355 360 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
370 375 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385 390 395 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
405 410 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
420 425 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
435 440 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
450 455 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465 470 475 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
485 490 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
500 505 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
515 520 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
530 535 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Gly Ala Ser Pro
545 550 555 560

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
565 570 575

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
580 585 590

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr

595 600 605

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
610 615 620

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
625 630 635 640

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
645 650 655

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
660 665 670

Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
675 680 685

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asp
690 695 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705 710 715 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
725 730 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
740 745 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
755 760 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
770 775 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785 790 795 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
805 810 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
820 825 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
835 840 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
850 855 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp

865 870 875 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
885 890 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
900 905 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
915 920 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
930 935 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945 950 955 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
965 970 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
980 985 990

His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val
995 1000 1005

Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr
1010 1015 1020

Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu
1025 1030 1035

Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His Gln
1040 1045 1050

Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
1055 1060 1065

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
1070 1075 1080

Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln
1085 1090 1095

Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp
1100 1105 1110

Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala
1115 1120 1125

Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu

```
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
```

1385 1390 1395

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1400 1405 1410

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1415 1420 1425

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1430 1435 1440

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1445 1450 1455

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1460 1465 1470

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1475 1480 1485

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1490 1495 1500

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1505 1510 1515

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1520 1525 1530

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1535 1540 1545

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1550 1555 1560

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1565 1570 1575

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1580 1585 1590

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1595 1600 1605

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1610 1615 1620

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1625 1630 1635

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly

1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 870
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 870
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ctcggtgagc tgcctgtgga cgcaagattt cctcctagag tgccaaaatc ttttccattc 120

ggcgcgccag gctccagccc ctccgcgagc acgggaaccg gaccaggttc gtcaccctca 180

gcatcaacgg ggacgggacc gggggcgtca ccaggaacgt cctccaccgg ctcgccgggt 240

gcatcacccg gaacgtcatc gaccggatcg ccagggagct cgacgccatc aggcgcaaca 300

ggatcacctg gctcaagccc tagcgcgtca accggcacgg gtccgggtgc ctcccctggc 360

acgtccagca ccggatcacc cggatcgagc ccatccgcct caaccggaac cggacccggt 420

acaccagggt cgggaacagc ctcctcgtca ccaggctcct caacccctc gggagccacg 480

ggttcgcccg gttcgtcaac gccttccgga gcaactggta gccccggagc atcgccagga 540

acttcgagca cggggtcgcc cgcctcgagc aacacctcag tcgtgtacaa aaagactctg 600

tttgtagaat tcacggatca ccttttcaac atcgctaagc caaggccacc ctggatgggt 660

ctgctaggtc ctaccatcca ggctgaggtt tatgatacag tggtcattac acttaagaac 720

atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag 780

ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct 840

ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

ggcctcattg gagccctact agtatgtaga gaagggagtc tggccaagga aaagacacag 1020

accttgcaca aatttatact acttttgct gtatttgatg aagggaaaag ttggcactca 1080

gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg 1200

aaatcagtct attggcatgt gattggaatg ggcaccactc ctgaagtgca ctcaatattc 1260

ctcgaaggtc acacatttct tgtgaggaac catcgccagg ctagcttgga aatctcgcca 1320

ataactttcc ttactgctca aacactcttg atggaccttg gacagtttct actgttttgt 1380

catatctctt cccaccaaca tgatggcatg gaagcttatg tcaaagtaga cagctgtcca 1440

gaggaacccc aactacgaat gaaaaataat gaagaagcgg aagactatga tgatgatctt 1500

actgattctg aaatggatgt ggtcaggttt gatgatgaca actctccttc ctttatccaa 1560

attcgctcag ttgccaagaa gcatcctaaa acttgggtac attacattgc tgctgaagag 1620

gaggactggg actatgctcc cttagtcctc gcccccgatg gcgcgccagg tgcctcgccg 1680

ggaacatcat caactggttc acccgggtca tccccctcgg cctcaaccgg gacgggtccc 1740

ggctcatccc ccagcgccag cactggaaca ggtcctggca ctcctggttc cggtacggca 1800

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actggttcgc cgggaactcc tggctcgggg acggcctcgt cgtcgcctgg ggcatcaccg 2040

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gcctcgagcg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 2160

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 2220

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 2280

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 2340

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 2400

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 2460

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 2520

agagatctag cttcaggact cattggccct ctcctcatct gctacaaaga atctgtagat 2580

caaagaggaa accagataat gtcagacaag aggaatgtca tcctgttttc tgtatttgat 2640

gagaaccgaa gctggtacct cacagagaat atacaacgct ttctccccaa tccagctgga 2700

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ggcatgaccg ccttactgaa ggtttctagt tgtgacaaga acactggtga ttattacgag 3060

gacagttatg aagatatttc agcatacttg ctgagtaaaa acaatgccat tgaaccaaga 3120

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gaagattttg acatttatga tgaggatgaa aatcagagcc cccgcagctt tcaaaagaaa 3300

acacgacact attttattgc tgcagtggag aggctctggg attatgggat gagtagctcc 3360

ccacatgttc taagaaacag ggctcagagt ggcagtgtcc ctcagttcaa gaaagttgtt 3420

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catttgggac tcctggggcc atatataaga gcagaagttg aagataatat catggtaact 3540

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gatcagaggc aaggagcaga acctagaaaa aactttgtca agcctaatga aaccaaaact 3660

tacttttgga aagtgcaaca tcatatggca cccactaaag atgagtttga ctgcaaagcc 3720

tgggcttatt tctctgatgt tgacctggaa aaagatgtgc actcaggcct gattggaccc 3780

cttctggtct gccacactaa cacactgaac cctgctcatg ggagacaagt gacagtacag 3840

gaatttgctc tgtttttcac catctttgat gagaccaaaa gctggtactt cactgaaaat 3900

atggaaagaa actgcagggc tccctgcaat atccagatgg aagatcccac ttttaaagag 3960

aattatcgct tccatgcaat caatggctac ataatggata cactacctgg cttagtaatg 4020

gctcaggatc aaaggattcg atggtatctg ctcagcatgg gcagcaatga aaacatccat 4080

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ctgtacaatc tctatccagg tgtttttgag acagtggaaa tgttaccatc caaagctgga 4200

atttggcggg tggaatgcct tattggcgag catctacatg ctgggatgag cacacttttt 4260

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tattccggat caatcaatgc ctggagcacc aaggagccct tttcttggat caaggtggat 4440

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agcctctaca tctctcagtt tatcatcatg tatagtcttg atgggaagaa gtggcagact 4560

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cattatagca ttcgcagcac tcttcgcatg gagttgatgg gctgtgattt aaatagttgc 4740

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tactttacca atatgtttgc cacctggtct ccttcaaaag ctcgacttca cctccaaggg 4860

aggagtaatg cctggagacc tcaggtgaat aatccaaaag agtggctgca agtggacttc 4920

cagaagacaa tgaaagtcac aggagtaact actcagggag taaaatctct gcttaccagc 4980

atgtatgtga aggagttcct catctccagc agtcaagatg gccatcagtg gactctcttt 5040

tttcagaatg gcaaagtaaa ggtttttcag ggaaatcaag actccttcac acctgtggtg 5100

aactctctag acccaccgtt actgactcgc taccttcgaa ttcaccccca gagttgggtg 5160

caccagattg ccctgaggat ggaggttctg ggctgcgagg cacaggacct ctacggcgcc 5220

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gcaaccggtc atcatcacca tcaccattga 5310

<210> 871
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 871
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Gly Ser Ser
20 25 30

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
35 40 45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
50 55 60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65 70 75 80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
85 90 95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
100 105 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
115 120 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
130 135 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145 150 155 160

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Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430
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Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Gly Ala Ser Pro
545                 550                 555                 560

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
                565                 570                 575

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            580                 585                 590

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
        595                 600                 605

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
    610                 615                 620

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
625                 630                 635                 640

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
                645                 650                 655

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
            660                 665                 670

Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
        675                 680                 685

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asp
    690                 695                 700
```

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Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975
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Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020

Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230
```

```
Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                 1240                 1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                 1255                 1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                 1270                 1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                 1285                 1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                 1300                 1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                 1315                 1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                 1330                 1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                 1345                 1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                 1360                 1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                 1375                 1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                 1390                 1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                 1405                 1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                 1420                 1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                 1435                 1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                 1450                 1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                 1465                 1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                 1480                 1485
```

```
Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1730                1735                1740
```

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 872
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 872
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ctcggtgagc tgcctgtggg cgcgccaggc tccagcccct ccgcgagcac gggaaccgga 120

ccaggttcgt caccctcagc atcaacgggg acgggaccgg gggcgtcacc aggaacgtcc 180

tccaccggct cgccgggtgc atcacccgga acgtcatcga ccggatcgcc agggagctcg 240

acgccatcag gcgcaacagg atcacctggc tcaagcccta gcgcgtcaac cggcacgggt 300

ccgggtgcct cccctggcac gtccagcacc ggatcacccg gatcgagccc atccgcctca 360

accggaaccg gacccggtac accagggtcg ggaacagcct cctcgtcacc aggctcctca 420

acccccctcgg gagccacggg ttcgcccggt tcgtcaacgc cttccggagc aactggtagc 480

cccggagcat cgccaggaac ttcgagcacg gggtcgcccg cctcgagcga cgcaagattt 540

cctcctagag tgccaaaatc ttttccattc aacacctcag tcgtgtacaa aaagactctg 600

tttgtagaat tcacggatca ccttttcaac atcgctaagc caaggccacc ctggatgggt 660

ctgctaggtc ctaccatcca ggctgaggtt tatgatacag tggtcattac acttaagaac 720

atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag 780

ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct 840

ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

ggcctcattg gagccctact agtatgtaga gaagggagtc tggccaagga aaagacacag 1020

accttgcaca aatttatact acttttgct gtatttgatg aagggaaaag ttggcactca 1080

gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg 1200

aaatcagtct attggcatgt gattggaatg ggcaccactc ctgaagtgca ctcaatattc 1260

ctcgaaggtc acacatttct tgtgaggaac catcgccagg ctagcttgga aatctcgcca 1320

ataactttcc ttactgctca aacactcttg atggaccttg gacagtttct actgttttgt 1380

catatctctt cccaccaaca tgatggcatg gaagcttatg tcaaagtaga cagctgtcca 1440

gaggaacccc aactacgaat gaaaaataat gaagaagcgg aagactatga tgatgatctt 1500

actgattctg aaatggatgt ggtcaggttt gatgatgaca actctccttc ctttatccaa 1560

attcgctcag ttgccaagaa gcatcctaaa acttgggtac attacattgc tgctgaagag 1620

gaggactggg actatgctcc cttagtcctc gcccccgatg gcgcgccagg tgcctcgccg 1680

ggaacatcat caactggttc acccgggtca tccccctcgg cctcaaccgg gacgggtccc 1740

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tcgtcatccc cgggaagctc aacaccgtcc ggagcgacag gatcacctgg ctcgtcacct 1860

tcggcgtcaa ctggaacggg gccaggggcc tcacccggaa cgtcctcgac tgggtcgcct 1920

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actggttcgc cgggaactcc tggctcgggg acggcctcgt cgtcgcctgg ggcatcaccg 2040

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gcctcgagcg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 2160

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 2220

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 2280

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 2340

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 2400

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 2460

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 2520

agagatctag cttcaggact cattggccct ctcctcatct gctacaaaga atctgtagat 2580

caaagaggaa accagataat gtcagacaag aggaatgtca tcctgttttc tgtatttgat 2640

gagaaccgaa gctggtacct cacagagaat atacaacgct ttctccccaa tccagctgga 2700

gtgcagcttg aggatccaga gttccaagcc tccaacatca tgcacagcat caatggctat 2760

gtttttgata gtttgcagtt gtcagtttgt ttgcatgagg tggcatactg gtacattcta 2820

agcattggag cacagactga cttccttttct gtcttcttct ctggatatac cttcaaacac 2880

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ggcatgaccg ccttactgaa ggtttctagt tgtgacaaga acactggtga ttattacgag 3060

gacagttatg aagatatttc agcatacttg ctgagtaaaa acaatgccat tgaaccaaga 3120

agcttctctc aaaacccacc agtcttgaaa cgccatcaac gggaaataac tcgtactact 3180

cttcagtcag atcaagagga aatcgattat gatgatacca tatcagttga aatgaagaag 3240

gaagattttg acatttatga tgaggatgaa aatcagagcc cccgcagctt tcaaaagaaa 3300

acacgacact attttattgc tgcagtggag aggctctggg attatgggat gagtagctcc 3360

ccacatgttc taagaaacag ggctcagagt ggcagtgtcc ctcagttcaa gaaagttgtt 3420

ttccaggaat ttactgatgg ctcctttact cagcccttat accgtggaga actaaatgaa 3480

catttgggac tcctggggcc atatataaga gcagaagttg aagataatat catggtaact 3540

ttcagaaatc aggcctctcg tccctattcc ttctattcta gccttatttc ttatgaggaa 3600

gatcagaggc aaggagcaga acctagaaaa aactttgtca agcctaatga aaccaaaact 3660

tacttttgga aagtgcaaca tcatatggca cccactaaag atgagtttga ctgcaaagcc 3720

tgggcttatt tctctgatgt tgacctggaa aaagatgtgc actcaggcct gattggaccc 3780

cttctggtct gccacactaa cacactgaac cctgctcatg ggagacaagt gacagtacag 3840

gaatttgctc tgtttttcac catctttgat gagaccaaaa gctggtactt cactgaaaat 3900

atggaaagaa actgcagggc tccctgcaat atccagatgg aagatcccac ttttaaagag 3960

aattatcgct tccatgcaat caatggctac ataatggata cactacctgg cttagtaatg 4020

gctcaggatc aaaggattcg atggtatctg ctcagcatgg gcagcaatga aaacatccat 4080

tctattcatt tcagtggaca tgtgttcact gtacgaaaaa aagaggagta taaaatggca 4140

ctgtacaatc tctatccagg tgtttttgag acagtggaaa tgttaccatc caaagctgga 4200

atttggcggg tggaatgcct tattggcgag catctacatg ctgggatgag cacacttttt 4260

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agcctctaca tctctcagtt tatcatcatg tatagtcttg atgggaagaa gtggcagact 4560

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ataaaacaca atatttttaa ccctccaatt attgctcgat acatccgttt gcacccaact 4680

cattatagca ttcgcagcac tcttcgcatg gagttgatgg gctgtgattt aaatagttgc 4740

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tactttacca atatgtttgc cacctggtct ccttcaaaag ctcgacttca cctccaaggg 4860

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cagaagacaa tgaaagtcac aggagtaact actcagggag taaaatctct gcttaccagc 4980

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aactctctag acccaccgtt actgactcgc taccttcgaa ttcaccccca gagttgggtg 5160

caccagattg ccctgaggat ggaggttctg ggctgcgagg cacaggacct ctacggcgcc 5220

ggatcacctg gggccgaaac ggccgaacaa aaactcatct cagaagagga tctgtcacct 5280

gcaaccggtc atcatcacca tcaccattga 5310

<210> 873
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 873

```
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1               5                   10                  15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Thr Ser Glu
            20                  25                  30

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
        35                  40                  45

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
    50                  55                  60

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
65                  70                  75                  80

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
                85                  90                  95

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
            100                 105                 110

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
        115                 120                 125

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
    130                 135                 140

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
145                 150                 155                 160

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ala Ser Ser
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255
```

```
Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525
```

```
Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Gly Ala Ser Pro
545                 550                 555                 560

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
                565                 570                 575

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            580                 585                 590

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
        595                 600                 605

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
    610                 615                 620

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
625                 630                 635                 640

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
                645                 650                 655

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
            660                 665                 670

Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
        675                 680                 685

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asp
    690                 695                 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800
```

```
Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020

Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065
```

```
Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320
```

```
Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575
```

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1580 1585 1590

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1595 1600 1605

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1610 1615 1620

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1625 1630 1635

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 874
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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gggagcgagc cagccacatc tgggtcggaa acgccaggca caagtgagtc tgcaactccc 180

gagtccggac ctggctccga gcctgccact agcggctccg agactccggg aacttccgag 240
agcgctacac cagaaagcgg acccggaacc agtaccgaac ctagcgaggg ctctgctccg 300
ggcagcccag ccggctctcc tacatccacg gaggagggca cttccgaatc cgccaccccg 360
gagtcagggc caggatctga acccgctacc tcaggcagtg agacgccagg aacgagcgag 420
tccgctacac cggagagtgg gccagggagc cctgctggat ctcctacgtc cactgaggaa 480
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cctcctagag tgccaaaatc ttttccattc aacacctcag tcgtgtacaa aaagactctg 600
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gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 2280

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 2340

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gacagttatg aagatatttc agcatacttg ctgagtaaaa acaatgccat tgaaccaaga 3120

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cttcagtcag atcaagagga aatcgattat gatgatacca tatcagttga aatgaagaag 3240

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acacgacact attttattgc tgcagtggag aggctctggg attatgggat gagtagctcc 3360

ccacatgttc taagaaacag ggctcagagt ggcagtgtcc ctcagttcaa gaaagttgtt 3420

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catttgggac tcctggggcc atatataaga gcagaagttg aagataatat catggtaact 3540

ttcagaaatc aggcctctcg tccctattcc ttctattcta gccttatttc ttatgaggaa 3600

gatcagaggc aaggagcaga acctagaaaa aactttgtca agcctaatga aaccaaaact 3660

tacttttgga aagtgcaaca tcatatggca cccactaaag atgagtttga ctgcaaagcc 3720

tgggcttatt tctctgatgt tgacctggaa aaagatgtgc actcaggcct gattggaccc 3780

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gaatttgctc tgtttttcac catctttgat gagaccaaaa gctggtactt cactgaaaat 3900

atggaaagaa actgcagggc tccctgcaat atccagatgg aagatcccac ttttaaagag 3960

aattatcgct tccatgcaat caatggctac ataatggata cactacctgg cttagtaatg 4020

gctcaggatc aaaggattcg atggtatctg ctcagcatgg gcagcaatga aaacatccat 4080

tctattcatt tcagtggaca tgtgttcact gtacgaaaaa aagaggagta taaaatggca 4140

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atttggcggg tggaatgcct tattggcgag catctacatg ctgggatgag cacacttttt 4260

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tatcgaggaa attccactgg aaccttaatg gtcttctttg gcaatgtgga ttcatctggg 4620

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tactttacca atatgtttgc cacctggtct ccttcaaaag ctcgacttca cctccaaggg 4860

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cagaagacaa tgaaagtcac aggagtaact actcagggag taaaatctct gcttaccagc 4980

atgtatgtga aggagttcct catctccagc agtcaagatg gccatcagtg gactctcttt 5040

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aactctctag acccaccgtt actgactcgc taccttcgaa ttcaccccca gagttgggtg 5160

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<210> 875
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 875
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Gly Ser Ser
20 25 30

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
35 40 45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
50 55 60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65 70 75 80

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Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
                85                  90                  95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
            100                 105                 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
        115                 120                 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
    130                 135                 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145                 150                 155                 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350
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Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620
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Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Ala Pro
            740                 745                 750

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
        755                 760                 765

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
    770                 775                 780

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
785                 790                 795                 800

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
                805                 810                 815

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
            820                 825                 830

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
        835                 840                 845

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
    850                 855                 860

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
865                 870                 875                 880

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                885                 890                 895
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Ala Ser Ser Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020

Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155
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Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410
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Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665
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Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 876
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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ccaggttcgt caccctcagc atcaacgggg acgggaccgg gggcgtcacc aggaacgtcc 180

tccaccggct cgccgggtgc atcacccgga acgtcatcga ccggatcgcc agggagctcg 240

acgccatcag gcgcaacagg atcacctggc tcaagcccta gcgcgtcaac cggcacgggt 300

ccgggtgcct cccctggcac gtccagcacc ggatcacccg gatcgagccc atccgcctca 360

accggaaccg gacccggtac accagggtcg ggaacagcct cctcgtcacc aggctcctca 420

acccccctcgg gagccacggg ttcgcccggt tcgtcaacgc cttccggagc aactggtagc 480

cccggagcat cgccaggaac ttcgagcacg gggtcgcccg cctcgagcga cgcaagattt 540

cctcctagag tgccaaaatc ttttccattc aacacctcag tcgtgtacaa aaagactctg 600

tttgtagaat tcacggatca ccttttcaac atcgctaagc caaggccacc ctggatgggt 660

ctgctaggtc ctaccatcca ggctgaggtt tatgatacag tggtcattac acttaagaac 720

atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag 780

ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct 840

ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

ggcctcattg gagccctact agtatgtaga gaagggagtc tggccaagga aaagacacag 1020

accttgcaca aatttatact acttttgct gtatttgatg aagggaaaag ttggcactca 1080

gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg 1200

aaatcagtct attggcatgt gattggaatg ggcaccactc ctgaagtgca ctcaatattc 1260

ctcgaaggtc acacatttct tgtgaggaac catcgccagg ctagcttgga aatctcgcca 1320

ataactttcc ttactgctca aacactcttg atggaccttg gacagtttct actgttttgt 1380

catatctctt cccaccaaca tgatggcatg gaagcttatg tcaaagtaga cagctgtcca 1440

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35 40 45

Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
50 55 60

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65 70 75 80

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro
85 90 95

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
100 105 110

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
115 120 125

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser
130 135 140

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145 150 155 160

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
165 170 175

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asn Thr

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Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
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Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
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Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
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His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
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Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
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Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
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Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Ala Pro
740 745 750

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
755 760 765

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
770 775 780

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
785 790 795 800

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
805 810 815

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
820 825 830

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
835 840 845

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
850 855 860

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
865 870 875 880

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
885 890 895

Ala Ser Ser Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
900 905 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
915 920 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
930 935 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945 950 955 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
965 970 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
980 985 990

His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val

995 1000 1005

Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr
1010 1015 1020

Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu
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Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His Gln
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Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
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Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
1070 1075 1080

Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln
1085 1090 1095

Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp
1100 1105 1110

Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala
1115 1120 1125

Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu
1130 1135 1140

Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
1145 1150 1155

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1160 1165 1170

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1175 1180 1185

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1190 1195 1200

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
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Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
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Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1235 1240 1245

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val

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Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
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Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
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Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
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Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
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Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
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Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
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Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
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Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
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Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
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Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
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Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
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Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
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Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
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Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
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Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
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Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
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Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1535 1540 1545

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
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Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1580 1585 1590

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1595 1600 1605

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1610 1615 1620

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1625 1630 1635

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
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Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
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Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
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Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
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aattatcgct tccatgcaat caatggctac ataatggata cactacctgg cttagtaatg 4020

gctcaggatc aaaggattcg atggtatctg ctcagcatgg gcagcaatga aaacatccat 4080

tctattcatt tcagtggaca tgtgttcact gtacgaaaaa aagaggagta taaaatggca 4140

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atttggcggg tggaatgcct tattggcgag catctacatg ctgggatgag cacacttttt 4260

ctggtgtaca gcaataagtg tcagactccc ctgggaatgg cttctggaca cattagagat 4320

tttcagatta cagcttcagg acaatatgga cagtgggccc caaagctggc cagacttcat 4380

tattccggat caatcaatgc ctggagcacc aaggagccct tttcttggat caaggtggat 4440

ctgttggcac caatgattat tcacggcatc aagacccagg gtgcccgtca gaagttctcc 4500

agcctctaca tctctcagtt tatcatcatg tatagtcttg atgggaagaa gtggcagact 4560

tatcgaggaa attccactgg aaccttaatg gtcttctttg gcaatgtgga ttcatctggg 4620

ataaaacaca atatttttaa ccctccaatt attgctcgat acatccgttt gcacccaact 4680

cattatagca ttcgcagcac tcttcgcatg gagttgatgg gctgtgattt aaatagttgc 4740

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tactttacca atatgtttgc cacctggtct ccttcaaaag ctcgacttca cctccaaggg 4860

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cagaagacaa tgaaagtcac aggagtaact actcagggag taaaatctct gcttaccagc 4980

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<210> 879
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 879
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Thr Ser Glu
            20                  25                  30

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
        35                  40                  45

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
    50                  55                  60

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
65                  70                  75                  80

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
                85                  90                  95

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
            100                 105                 110

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
        115                 120                 125

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
    130                 135                 140

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
145                 150                 155                 160

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ala Ser Ser
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285
```

```
Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560
```

```
Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Ala Pro
            740                 745                 750

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
        755                 760                 765

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
    770                 775                 780

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
785                 790                 795                 800

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
                805                 810                 815

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
            820                 825                 830
```

```
Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
        835                 840                 845

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
    850                 855                 860

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
865                 870                 875                 880

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                885                 890                 895

Ala Ser Ser Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020

Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095
```

```
Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350
```

```
Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                 1360                 1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                 1375                 1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                 1390                 1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                 1405                 1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                 1420                 1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                 1435                 1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                 1450                 1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                 1465                 1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                 1480                 1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                 1495                 1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                 1510                 1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                 1525                 1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                 1540                 1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                 1555                 1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                 1570                 1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                 1585                 1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                 1600                 1605
```

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1610 1615 1620

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1625 1630 1635

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 880
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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gggagcgagc cagccacatc tgggtcggaa acgccaggca caagtgagtc tgcaactccc 180

gagtccggac ctggctccga gcctgccact agcggctccg agactccggg aacttccgag 240

agcgctacac cagaaagcgg acccggaacc agtaccgaac ctagcgaggg ctctgctccg 300

ggcagcccag ccggctctcc tacatccacg gaggagggca cttccgaatc cgccaccccg 360

gagtcagggc caggatctga acccgctacc tcaggcagtg agacgccagg aacgagcgag 420

tccgctacac cggagagtgg gccagggagc cctgctggat ctcctacgtc cactgaggaa 480

gggtcaccag cgggctcgcc caccagcact gaagaaggtg cctcgagcga cgcaagattt 540

cctcctagag tgccaaaatc ttttccattc aacacctcag tcgtgtacaa aaagactctg 600

tttgtagaat tcacggatca ccttttcaac atcgctaagc caaggccacc ctggatgggt 660

ctgctaggtc ctaccatcca ggctgaggtt tatgatacag tggtcattac acttaagaac 720

atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag 780

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ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

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atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg 1200

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<210> 881
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 881
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Gly Ala Pro Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
20 25 30

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
35 40 45

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
50 55 60

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
65 70 75 80

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
85 90 95

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
100 105 110

```
Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
        115                 120                 125

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
    130                 135                 140

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
145                 150                 155                 160

Ser Thr Glu Glu Gly Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380
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Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655
```

```
Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Ala Pro
            740                 745                 750

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
        755                 760                 765

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
    770                 775                 780

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
785                 790                 795                 800

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
                805                 810                 815

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
            820                 825                 830

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
        835                 840                 845

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
    850                 855                 860

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
865                 870                 875                 880

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                885                 890                 895

Ala Ser Ser Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925
```

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Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020

Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185
```

```
Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                 1195                 1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                 1210                 1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                 1225                 1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                 1240                 1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                 1255                 1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                 1270                 1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                 1285                 1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                 1300                 1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                 1315                 1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                 1330                 1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                 1345                 1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                 1360                 1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                 1375                 1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                 1390                 1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                 1405                 1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                 1420                 1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                 1435                 1440
```

```
Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695
```

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 882
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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tcggaaacgc caggcacaag tgagtctgca actcccgagt ccggacctgg ctccgagcct 180

gccactagcg gctccgagac tccgggaact tccgagagcg ctacaccaga aagcggaccc 240

ggaaccagta ccgaacctag cgagggctct gctccgggca gcccagccgg ctctcctaca 300

tccacggagg agggcacttc cgaatccgcc accccggagt cagggccagg atctgaaccc 360

gctacctcag gcagtgagac gccaggaacg agcgagtccg ctacaccgga gagtgggcca 420

gggagccctg ctggatctcc tacgtccact gaggaagggt caccagcggg ctcgcccacc 480

agcactgaag aaggtgcctc gagcagtgat ctcggtgagc tgcctgtgga cgcaagattt 540

cctcctagag tgccaaaatc ttttccattc aacacctcag tcgtgtacaa aaagactctg 600

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ctgctaggtc ctaccatcca ggctgaggtt tatgatacag tggtcattac acttaagaac 720

atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag 780

ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct 840

ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

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gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

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cctttacttt atggggaagt tggagacaca ctgttgatta tatttaagaa tcaagcaagc 1860

agaccatata acatctaccc tcacggaatc actgatgtcc gtcctttgta ttcaaggaga 1920

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<210> 883
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 883
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1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Gly Ala Pro Thr Ser Glu Ser Ala
35 40 45

Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu
50 55 60

Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
65 70 75 80

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala
85 90 95

Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
100 105 110

Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr
115 120 125

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr
130 135 140

Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
145 150 155 160

Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser
165 170 175

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ala Ser Ser Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

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Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480
```

```
Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Ala Pro
            740                 745                 750
```

```
Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
        755                 760                 765

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
    770                 775                 780

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
785                 790                 795                 800

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
                805                 810                 815

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
            820                 825                 830

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
        835                 840                 845

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
    850                 855                 860

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
865                 870                 875                 880

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                885                 890                 895

Ala Ser Ser Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
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```

```
Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
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Pro Arg  Ser Phe Ser Gln Asn  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275
```

```
Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530
```

```
Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1730                1735                1740

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
    1745                1750                1755

Ser Pro  Ala Thr Gly His His  His His His His
    1760                1765
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<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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tctgggtcgg aaacgccagg cacaagtgag tctgcaactc ccgagtccgg acctggctcc 240

gagcctgcca ctagcggctc cgagactccg ggaacttccg agagcgctac accagaaagc 300

ggacccggaa ccagtaccga acctagcgag ggctctgctc cgggcagccc agccggctct 360

cctacatcca cggaggaggg cacttccgaa tccgccaccc cggagtcagg gccaggatct 420

gaacccgcta cctcaggcag tgagacgcca ggaacgagcg agtccgctac accggagagt 480

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gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

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<210> 885
<211> 1759
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 885
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys

35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met

305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln

580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
740 745 750

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
755 760 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
770 775 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785 790 795 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
805 810 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
820 825 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
835 840 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly

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    850                 855                 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865                 870                 875                 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
```

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Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1130 1135 1140

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1145 1150 1155

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1160 1165 1170

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1175 1180 1185

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1190 1195 1200

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1205 1210 1215

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1220 1225 1230

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1235 1240 1245

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1250 1255 1260

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1265 1270 1275

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1280 1285 1290

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
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Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1310 1315 1320

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1325 1330 1335

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1340 1345 1350

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1355 1360 1365

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe

1370 1375 1380

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1385 1390 1395

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1400 1405 1410

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1415 1420 1425

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1430 1435 1440

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1445 1450 1455

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1460 1465 1470

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1475 1480 1485

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1490 1495 1500

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1505 1510 1515

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1520 1525 1530

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Pro Pro Gly
1580 1585 1590

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
1595 1600 1605

Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr
1610 1615 1620

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly

1625 1630 1635

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
1640 1645 1650

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
1655 1660 1665

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
1670 1675 1680

Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
1685 1690 1695

Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
1700 1705 1710

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
1715 1720 1725

Ser Gly Thr Ala Ser Ser Ser Ala Ser Ser Glu Gln Lys Leu Ile
1730 1735 1740

Ser Glu Glu Asp Leu Ser Pro Ala Thr Gly His His His His His
1745 1750 1755

His

<210> 886
<211> 5280
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

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gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720
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cgccatcaac gggaaataac tcgtactact cttcagtcag atcaagagga aatcgattat 2760

gatgatacca tatcagttga aatgaagaag gaagattttg acatttatga tgaggatgaa 2820

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aggctctggg attatgggat gagtagctcc ccacatgttc taagaaacag ggctcagagt 2940

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cccactaaag atgagtttga ctgcaaagcc tgggcttatt tctctgatgt tgacctggaa 3300

aaagatgtgc actcaggcct gattggaccc cttctggtct gccacactaa cacactgaac 3360

cctgctcatg ggagacaagt gacagtacag gaatttgctc tgtttttcac catctttgat 3420

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<210> 887
<211> 1903
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

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Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415
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Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685
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Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Pro Gly Ala Ser
            740                 745                 750

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
        755                 760                 765

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
    770                 775                 780

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro
785                 790                 795                 800

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
                805                 810                 815

Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
            820                 825                 830

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
        835                 840                 845

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
    850                 855                 860

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
865                 870                 875                 880

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
                885                 890                 895

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
            900                 905                 910

Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
        915                 920                 925

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
    930                 935                 940

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
945                 950                 955                 960
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Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
               965                 970                 975

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
           980                 985                 990

Pro Gly Thr Ser Ser Thr Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser
       995                 1000                1005

Thr Gly  Thr Gly Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
   1010                1015                1020

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Ala Ser
   1025                1030                1035

Ser Pro  Pro Val Leu Lys Arg  His Gln Ala Glu Ile  Thr Arg Thr
   1040                1045                1050

Thr Leu  Gln Ser Asp Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile
   1055                1060                1065

Ser Val  Glu Met Lys Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp
   1070                1075                1080

Glu Asn  Gln Ser Pro Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr
   1085                1090                1095

Phe Ile  Ala Ala Val Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser
   1100                1105                1110

Ser Pro  His Val Leu Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro
   1115                1120                1125

Gln Phe  Lys Lys Val Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe
   1130                1135                1140

Thr Gln  Pro Leu Tyr Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu
   1145                1150                1155

Leu Gly  Pro Tyr Ile Arg Ala  Glu Val Glu Asp Asn  Ile Met Val
   1160                1165                1170

Thr Phe  Arg Asn Gln Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser
   1175                1180                1185

Leu Ile  Ser Tyr Glu Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg
   1190                1195                1200

Lys Asn  Phe Val Lys Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys
   1205                1210                1215
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Val Gln  His His Met Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys
    1220                 1225                 1230

Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu Glu Lys  Asp Val His
    1235                 1240                 1245

Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys His Thr  Asn Thr Leu
    1250                 1255                 1260

Asn Pro  Ala His Gly Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu
    1265                 1270                 1275

Phe Phe  Thr Ile Phe Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu
    1280                 1285                 1290

Asn Met  Glu Arg Asn Cys Arg  Ala Pro Cys Asn Ile  Gln Met Glu
    1295                 1300                 1305

Asp Pro  Thr Phe Lys Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly
    1310                 1315                 1320

Tyr Ile  Met Asp Thr Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln
    1325                 1330                 1335

Arg Ile  Arg Trp Tyr Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile
    1340                 1345                 1350

His Ser  Ile His Phe Ser Gly  His Val Phe Thr Val  Arg Lys Lys
    1355                 1360                 1365

Glu Glu  Tyr Lys Met Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe
    1370                 1375                 1380

Glu Thr  Val Glu Met Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val
    1385                 1390                 1395

Glu Cys  Leu Ile Gly Glu His  Leu His Ala Gly Met  Ser Thr Leu
    1400                 1405                 1410

Phe Leu  Val Tyr Ser Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala
    1415                 1420                 1425

Ser Gly  His Ile Arg Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr
    1430                 1435                 1440

Gly Gln  Trp Ala Pro Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser
    1445                 1450                 1455

Ile Asn  Ala Trp Ser Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val
    1460                 1465                 1470
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Asp Leu  Leu Ala Pro Met Ile  Ile His Gly Ile Lys  Thr Gln Gly
    1475                1480                1485

Ala Arg  Gln Lys Phe Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile
    1490                1495                1500

Met Tyr  Ser Leu Asp Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn
    1505                1510                1515

Ser Thr  Gly Thr Leu Met Val  Phe Phe Gly Asn Val  Asp Ser Ser
    1520                1525                1530

Gly Ile  Lys His Asn Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr
    1535                1540                1545

Ile Arg  Leu His Pro Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg
    1550                1555                1560

Met Glu  Leu Met Gly Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu
    1565                1570                1575

Gly Met  Glu Ser Lys Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser
    1580                1585                1590

Ser Tyr  Phe Thr Asn Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala
    1595                1600                1605

Arg Leu  His Leu Gln Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val
    1610                1615                1620

Asn Asn  Pro Lys Glu Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met
    1625                1630                1635

Lys Val  Thr Gly Val Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr
    1640                1645                1650

Ser Met  Tyr Val Lys Glu Phe  Leu Ile Ser Ser Ser  Gln Asp Gly
    1655                1660                1665

His Gln  Trp Thr Leu Phe Phe  Gln Asn Gly Lys Val  Lys Val Phe
    1670                1675                1680

Gln Gly  Asn Gln Asp Ser Phe  Thr Pro Val Val Asn  Ser Leu Asp
    1685                1690                1695

Pro Pro  Leu Leu Thr Arg Tyr  Leu Arg Ile His Pro  Gln Ser Trp
    1700                1705                1710

Val His  Gln Ile Ala Leu Arg  Met Glu Val Leu Gly  Cys Glu Ala
    1715                1720                1725
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Gln Asp Leu Tyr Gly Ala Pro Pro Gly Ser Ser Pro Ser Ala Ser
1730 1735 1740

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr
1745 1750 1755

Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
1760 1765 1770

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
1775 1780 1785

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser
1790 1795 1800

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
1805 1810 1815

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
1820 1825 1830

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
1835 1840 1845

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
1850 1855 1860

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
1865 1870 1875

Ser Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
1880 1885 1890

Pro Ala Thr Gly His His His His His His
1895 1900

<210> 888
<211> 5712
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

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20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

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Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320
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Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590
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Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
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Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Thr Pro Gly
            740                 745                 750

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
        755                 760                 765

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    770                 775                 780

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
785                 790                 795                 800

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
                805                 810                 815

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            820                 825                 830

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
        835                 840                 845

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
    850                 855                 860
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Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
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Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125
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Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
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Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
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Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380
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Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
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Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
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Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
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Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                1465                1470

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                1480                1485

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1490                1495                1500

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                1510                1515

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1520                1525                1530

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1535                1540                1545

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1550                1555                1560

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
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Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
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Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
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Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
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Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
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Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
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Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
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Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1775                 1780                 1785

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
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Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
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Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
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Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Glu Pro Ala
    1835                 1840                 1845

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1850                 1855                 1860

Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    1865                 1870                 1875

Pro Ala  Ser Ser Glu Gln Lys  Leu Ile Ser Glu Glu  Asp Leu Ser
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

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Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
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Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495
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His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
            740                 745                 750

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
        755                 760                 765
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Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
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Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
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Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                805                 810                 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            820                 825                 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
        835                 840                 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    850                 855                 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865                 870                 875                 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
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Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
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Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
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Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
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Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
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Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
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Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
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Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
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Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
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Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
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Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
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Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
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Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
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Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
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Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
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Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
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Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
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Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
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Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
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Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
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Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
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Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
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Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
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Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
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Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                1510                1515

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
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Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
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Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
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Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
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Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Pro Gly
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Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser Pro Gly  Ala Ser Pro
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Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Thr Pro Gly  Ser Gly Thr
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Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly
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Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Thr Pro Gly
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Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser Ser Thr  Pro Ser Gly
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Ala Thr  Gly Ser Pro Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly
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Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr Gly Thr  Gly Pro Gly
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Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly Pro Gly  Ala Ser Pro
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Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Thr Pro Gly  Ser Gly Thr
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Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly
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Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr Gly Thr  Gly Pro Gly
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Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly Pro Gly  Ala Ser Pro
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Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
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Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
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Pro Ala Thr Gly His His His His His His
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
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Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val

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Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
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Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
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Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro

660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Thr Ser Glu
740 745 750

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
755 760 765

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
770 775 780

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
785 790 795 800

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835 840 845

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
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Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
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Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
915 920 925

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930 935 940

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
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Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
965 970 975

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
980 985 990

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
995 1000 1005

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1010 1015 1020

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Ala Ser
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Ser Pro Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr
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Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
1055 1060 1065

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp
1070 1075 1080

Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr
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Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser
1100 1105 1110

Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro
1115 1120 1125

Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe
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Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
1145 1150 1155

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val
1160 1165 1170

Thr Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser
1175 1180 1185

Leu Ile Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg

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Lys Asn  Phe Val Lys Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys
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Val Gln  His His Met Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys
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Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu Glu Lys  Asp Val His
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Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys His Thr  Asn Thr Leu
    1250                1255                1260

Asn Pro  Ala His Gly Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu
    1265                1270                1275

Phe Phe  Thr Ile Phe Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu
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Asn Met  Glu Arg Asn Cys Arg  Ala Pro Cys Asn Ile  Gln Met Glu
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Asp Pro  Thr Phe Lys Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly
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Tyr Ile  Met Asp Thr Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln
    1325                1330                1335

Arg Ile  Arg Trp Tyr Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile
    1340                1345                1350

His Ser  Ile His Phe Ser Gly  His Val Phe Thr Val  Arg Lys Lys
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Glu Glu  Tyr Lys Met Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe
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Glu Thr  Val Glu Met Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val
    1385                1390                1395

Glu Cys  Leu Ile Gly Glu His  Leu His Ala Gly Met  Ser Thr Leu
    1400                1405                1410

Phe Leu  Val Tyr Ser Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala
    1415                1420                1425

Ser Gly  His Ile Arg Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr
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Gly Gln  Trp Ala Pro Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser
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    1475                1480                1485

Ala Arg  Gln Lys Phe Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile
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Met Tyr  Ser Leu Asp Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn
    1505                1510                1515

Ser Thr  Gly Thr Leu Met Val  Phe Phe Gly Asn Val  Asp Ser Ser
    1520                1525                1530

Gly Ile  Lys His Asn Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr
    1535                1540                1545

Ile Arg  Leu His Pro Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg
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Met Glu  Leu Met Gly Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu
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Gly Met  Glu Ser Lys Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser
    1580                1585                1590

Ser Tyr  Phe Thr Asn Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala
    1595                1600                1605

Arg Leu  His Leu Gln Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val
    1610                1615                1620

Asn Asn  Pro Lys Glu Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met
    1625                1630                1635

Lys Val  Thr Gly Val Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr
    1640                1645                1650

Ser Met  Tyr Val Lys Glu Phe  Leu Ile Ser Ser Ser  Gln Asp Gly
    1655                1660                1665

His Gln  Trp Thr Leu Phe Phe  Gln Asn Gly Lys Val  Lys Val Phe
    1670                1675                1680

Gln Gly  Asn Gln Asp Ser Phe  Thr Pro Val Val Asn  Ser Leu Asp
    1685                1690                1695

Pro Pro  Leu Leu Thr Arg Tyr  Leu Arg Ile His Pro  Gln Ser Trp
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Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly Cys Glu Ala
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1745 1750 1755

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1835 1840 1845

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1880 1885 1890

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser
1895 1900 1905

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
1910 1915 1920

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1925 1930 1935

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser
1940 1945 1950

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu

1955 1960 1965

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
1970 1975 1980

Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1985 1990 1995

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
2000 2005 2010

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Ala Ser Ser Glu Gln
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Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala Thr Gly His His
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aaccctccaa ttattgctcg atacatccgt ttgcacccaa ctcattatag cattcgcagc 4680

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<210> 895
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 895
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Gly Ser Ser
20 25 30

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
35 40 45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
50 55 60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65 70 75 80

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Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
                85                  90                  95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
            100                 105                 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
        115                 120                 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
    130                 135                 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145                 150                 155                 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350
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Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Gly Ala Ser Pro
545                 550                 555                 560

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
                565                 570                 575

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            580                 585                 590

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
        595                 600                 605

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
    610                 615                 620
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Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
625                 630                 635                 640

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
                645                 650                 655

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
            660                 665                 670

Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
        675                 680                 685

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asp
    690                 695                 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895
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Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020

Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155
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Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                 1165                 1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                 1180                 1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                 1195                 1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                 1210                 1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                 1225                 1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                 1240                 1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                 1255                 1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                 1270                 1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                 1285                 1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                 1300                 1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                 1315                 1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                 1330                 1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                 1345                 1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                 1360                 1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                 1375                 1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                 1390                 1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                 1405                 1410
```

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Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665
```

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
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Ser Pro Ala Thr Gly His His His His His His
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<210> 896
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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<210> 897
<211> 1772
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 897
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

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Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445
```

```
Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720
```

```
Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Gly Ala Pro
            820                 825                 830

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
        835                 840                 845

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
    850                 855                 860

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
865                 870                 875                 880

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
                885                 890                 895

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
            900                 905                 910

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
        915                 920                 925

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
    930                 935                 940

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
945                 950                 955                 960

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                965                 970                 975

Ala Ser Ser Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val
            980                 985                 990
```

```
Val Phe Gln Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
        995                 1000                1005

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1010                1015                1020

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1025                1030                1035

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1040                1045                1050

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1055                1060                1065

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1070                1075                1080

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1085                1090                1095

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1100                1105                1110

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1115                1120                1125

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1130                1135                1140

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1145                1150                1155

Arg Gly  Ala Pro Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1160                1165                1170

Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser
    1175                1180                1185

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
    1190                1195                1200

Ser Gly  Ser Glu Thr Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1205                1210                1215

Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1220                1225                1230

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Pro
    1235                1240                1245
```

```
Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Ser Pro  Ala Gly Ser
    1250                 1255                 1260

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1265                 1270                 1275

Thr Glu  Glu Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1280                 1285                 1290

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Ala Ser
    1295                 1300                 1305

Ser Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1310                 1315                 1320

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1325                 1330                 1335

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1340                 1345                 1350

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1355                 1360                 1365

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1370                 1375                 1380

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1385                 1390                 1395

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1400                 1405                 1410

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1415                 1420                 1425

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1430                 1435                 1440

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1445                 1450                 1455

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1460                 1465                 1470

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1475                 1480                 1485

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1490                 1495                 1500
```

```
Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1505                1510                1515

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1520                1525                1530

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1535                1540                1545

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1550                1555                1560

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1565                1570                1575

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1580                1585                1590

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1595                1600                1605

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1610                1615                1620

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1625                1630                1635

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1640                1645                1650

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1655                1660                1665

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1670                1675                1680

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1685                1690                1695

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    1700                1705                1710

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    1715                1720                1725

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    1730                1735                1740

Gly Ser  Pro Gly Ala Glu Thr  Ala Glu Gln Lys Leu  Ile Ser Glu
    1745                1750                1755
```

Glu Asp Leu Ser Pro Ala Thr Gly His His His His His His
1760 1765 1770

<210> 898
<211> 5319
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

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attagagatt ttcagattac agcttcagga caatatggac agtgggcccc aaagctggcc 4380

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agttgggtgc accagattgc cctgaggatg gaggttctgg gctgcgaggc acaggacctc 5220

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ctgtcacctg caaccggtca tcatcaccat caccattga 5319

<210> 899
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 899

```
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1               5                   10                  15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270
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```
Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540
```

```
Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815
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Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Gly
            820                 825                 830

Ala Pro Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
        835                 840                 845

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
    850                 855                 860

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
865                 870                 875                 880

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
                885                 890                 895

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
            900                 905                 910

Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
        915                 920                 925

Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro
    930                 935                 940

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr
945                 950                 955                 960

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
                965                 970                 975

Pro Gly Ala Ser Ser Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080
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```
Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1145                1150                1155

Pro Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    1160                1165                1170

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1175                1180                1185

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1190                1195                1200

Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1205                1210                1215

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1220                1225                1230

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Pro  Ala Gly Ser
    1235                1240                1245

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1250                1255                1260

Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1265                1270                1275

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1280                1285                1290

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ala Ser  Ser Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335
```

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Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590
```

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1595 1600 1605

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1610 1615 1620

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1625 1630 1635

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 900
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 900
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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

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gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

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gcaaccggtc atcatcacca tcaccattga 5310

<210> 901
<211> 1772
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 901
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala

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            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
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370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr

645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
740 745 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
755 760 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
770 775 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785 790 795 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
805 810 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Gly Ala Pro
820 825 830

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
835 840 845

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
850 855 860

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
865 870 875 880

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
885 890 895

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
900 905 910

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro

915 920 925

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
930 935 940

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
945 950 955 960

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
965 970 975

Ala Ser Ser Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val
980 985 990

Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg
995 1000 1005

Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
1010 1015 1020

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala
1025 1030 1035

Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu
1040 1045 1050

Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro
1055 1060 1065

Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala
1070 1075 1080

Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser
1085 1090 1095

Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
1100 1105 1110

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg
1115 1120 1125

Gln Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp
1130 1135 1140

Glu Thr Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys
1145 1150 1155

Arg Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
1160 1165 1170

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala

1175 1180 1185

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
1190 1195 1200

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
1205 1210 1215

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
1220 1225 1230

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
1235 1240 1245

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
1250 1255 1260

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
1265 1270 1275

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
1280 1285 1290

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser
1295 1300 1305

Ser Ala Pro Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu
1310 1315 1320

Asn Tyr Arg Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu
1325 1330 1335

Pro Gly Leu Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu
1340 1345 1350

Leu Ser Met Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser
1355 1360 1365

Gly His Val Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala
1370 1375 1380

Leu Tyr Asn Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu
1385 1390 1395

Pro Ser Lys Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu
1400 1405 1410

His Leu His Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn
1415 1420 1425

Lys Cys Gln Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp

1430 1435 1440

Phe Gln Ile Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys
1445 1450 1455

Leu Ala Arg Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr
1460 1465 1470

Lys Glu Pro Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met
1475 1480 1485

Ile Ile His Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser
1490 1495 1500

Ser Leu Tyr Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly
1505 1510 1515

Lys Lys Trp Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met
1520 1525 1530

Val Phe Phe Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile
1535 1540 1545

Phe Asn Pro Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr
1550 1555 1560

His Tyr Ser Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys
1565 1570 1575

Asp Leu Asn Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala
1580 1585 1590

Ile Ser Asp Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met
1595 1600 1605

Phe Ala Thr Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly
1610 1615 1620

Arg Ser Asn Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp
1625 1630 1635

Leu Gln Val Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr
1640 1645 1650

Thr Gln Gly Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu
1655 1660 1665

Phe Leu Ile Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe
1670 1675 1680

Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser

1685 1690 1695

Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg
1700 1705 1710

Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu
1715 1720 1725

Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala
1730 1735 1740

Gly Ser Pro Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu
1745 1750 1755

Glu Asp Leu Ser Pro Ala Thr Gly His His His His His His
1760 1765 1770

<210> 902
<211> 5319
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

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caaagaggaa accagataat gtcagacaag aggaatgtca tcctgttttc tgtatttgat 1740

gagaaccgaa gctggtacct cacagagaat atacaacgct ttctccccaa tccagctgga 1800

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gtttttgata gtttgcagtt gtcagtttgt ttgcatgagg tggcatactg gtacattcta 1920

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cttcagtcag atcaagagga aatcgattat gatgatacca tatcagttga aatgaagaag 2340

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gctcagagtg gcagtgtccc tcagttcaag aaagttgttt tccaggaatt tactgatggc 3000

tcctttactc agcccttata ccgtggagaa ctaaatgaac atttgggact cctggggcca 3060

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ctgtcacctg caaccggtca tcatcaccat caccattga 5319

<210> 903
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 903
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

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His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480
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Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750
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Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Gly Ala Pro Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
            820                 825                 830

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
        835                 840                 845

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
    850                 855                 860

Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
865                 870                 875                 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
                885                 890                 895

Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
            900                 905                 910

Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
        915                 920                 925

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
    930                 935                 940

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
945                 950                 955                 960

Ser Ala Pro Gly Ala Ser Ser Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020
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Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gly Ala Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    1160                1165                1170

Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1175                1180                1185

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1190                1195                1200

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala
    1205                1210                1215

Ser Thr  Gly Thr Gly Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    1220                1225                1230

Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1235                1240                1245

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1250                1255                1260

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser
    1265                1270                1275
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Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    1280                 1285                 1290

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1295                 1300                 1305

Ala Ser  Ser Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                 1315                 1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                 1330                 1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                 1345                 1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                 1360                 1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                 1375                 1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                 1390                 1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                 1405                 1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                 1420                 1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                 1435                 1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                 1450                 1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                 1465                 1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                 1480                 1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                 1495                 1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                 1510                 1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                 1525                 1530
```

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Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1730                1735                1740

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
    1745                1750                1755

Ser Pro  Ala Thr Gly His His  His His His His His
    1760                1765
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<210> 904
<211> 5310

<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

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<211> 1772
<212> PRT
<213> Artificial Sequence

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Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300
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Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
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Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575
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Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Gly Ala Pro
            820                 825                 830

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
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Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
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Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
865                 870                 875                 880

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
                885                 890                 895

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
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Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
        915                 920                 925

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
    930                 935                 940

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
945                 950                 955                 960

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                965                 970                 975

Ala Ser Ser Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val
            980                 985                 990

Val Phe Gln Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
        995                 1000                1005

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1010                1015                1020

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1025                1030                1035

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1040                1045                1050

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1055                1060                1065

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1070                1075                1080

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1085                1090                1095

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1100                1105                1110
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Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
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Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1130                1135                1140

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1145                1150                1155

Arg Ala  Pro Cys Asn Ile Gly  Ala Pro Gly Thr Pro  Gly Ser Gly
    1160                1165                1170

Thr Ala  Ser Ser Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
    1175                1180                1185

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1190                1195                1200

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser
    1205                1210                1215

Pro Ser  Ala Ser Thr Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly
    1220                1225                1230

Thr Ala  Ser Ser Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
    1235                1240                1245

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1250                1255                1260

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser
    1265                1270                1275

Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser
    1280                1285                1290

Gly Ala  Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
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Gly Ser  Pro Ala Ser Ser Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1310                1315                1320

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1325                1330                1335

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1340                1345                1350

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1355                1360                1365
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Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
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Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1385                1390                1395

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1400                1405                1410

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1415                1420                1425

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1430                1435                1440

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1445                1450                1455

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1460                1465                1470

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1475                1480                1485

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1490                1495                1500

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1505                1510                1515

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1520                1525                1530

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1535                1540                1545

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1550                1555                1560

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1565                1570                1575

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1580                1585                1590

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1595                1600                1605

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1610                1615                1620
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Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser
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Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg
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Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu
1715 1720 1725

Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala
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<211> 1769
<212> PRT
<213> Artificial Sequence

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20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

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Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
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Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400
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Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
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Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670
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Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Gly
            820                 825                 830

Ala Pro Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
        835                 840                 845

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
    850                 855                 860

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
865                 870                 875                 880

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
                885                 890                 895

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
            900                 905                 910

Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
        915                 920                 925

Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro
    930                 935                 940
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Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr
945                 950                 955                 960

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
                965                 970                 975

Pro Gly Ala Ser Ser Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gly Ala Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    1160                1165                1170

Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1175                1180                1185

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1190                1195                1200
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Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala
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Ser Thr  Gly Thr Gly Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
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Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1235                 1240                 1245

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1250                 1255                 1260

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser
    1265                 1270                 1275

Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    1280                 1285                 1290

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1295                 1300                 1305

Ala Ser  Ser Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                 1315                 1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                 1330                 1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                 1345                 1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                 1360                 1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                 1375                 1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                 1390                 1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                 1405                 1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                 1420                 1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                 1435                 1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                 1450                 1455
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Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710
```

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
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Ser Pro Ala Thr Gly His His His His His His
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tatcgaggaa attccactgg aaccttaatg gtcttctttg gcaatgtgga ttcatctggg 4620

ataaaacaca atatttttaa ccctccaatt attgctcgat acatccgttt gcacccaact 4680

cattatagca ttcgcagcac tcttcgcatg gagttgatgg gctgtgattt aaatagttgc 4740

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tactttacca atatgtttgc cacctggtct ccttcaaaag ctcgacttca cctccaaggg 4860

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cagaagacaa tgaaagtcac aggagtaact actcagggag taaaatctct gcttaccagc 4980

atgtatgtga aggagttcct catctccagc agtcaagatg gccatcagtg gactctcttt 5040

tttcagaatg gcaaagtaaa ggtttttcag ggaaatcaag actccttcac acctgtggtg 5100

aactctctag acccaccgtt actgactcgc taccttcgaa ttcaccccca gagttgggtg 5160

caccagattg ccctgaggat ggaggttctg ggctgcgagg cacaggacct ctacggcgcc 5220

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gcaaccggtc atcatcacca tcaccattga 5310

<210> 909
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 909
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Gly Ser Ser
20 25 30

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
35 40 45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
50 55 60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65 70 75 80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
85 90 95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
100 105 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
115 120 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
130 135 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145 150 155 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn

225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
260 265 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
275 280 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
290 295 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305 310 315 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
325 330 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
340 345 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
355 360 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
370 375 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385 390 395 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
405 410 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
420 425 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
435 440 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
450 455 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465 470 475 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
485 490 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp

500 505 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
515 520 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
530 535 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545 550 555 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
565 570 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
580 585 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
595 600 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
610 615 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
645 650 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val

770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
835 840 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
885 890 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
900 905 910

Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
915 920 925

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
930 935 940

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
945 950 955 960

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
965 970 975

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly
980 985 990

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
995 1000 1005

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
1010 1015 1020

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
1025 1030 1035

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr

1040 1045 1050

Ser Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Gln Glu Glu Ile
1055 1060 1065

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
1070 1075 1080

Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln
1085 1090 1095

Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp
1100 1105 1110

Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala
1115 1120 1125

Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu
1130 1135 1140

Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
1145 1150 1155

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1160 1165 1170

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1175 1180 1185

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1190 1195 1200

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1205 1210 1215

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1220 1225 1230

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1235 1240 1245

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1250 1255 1260

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1265 1270 1275

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1280 1285 1290

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro

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    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
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Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
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1550 1555 1560

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1565 1570 1575

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1580 1585 1590

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1595 1600 1605

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1610 1615 1620

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1625 1630 1635

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 910
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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ccaggttcgt caccctcagc atcaacgggg acgggaccgg gggcgtcacc aggaacgtcc 180

tccaccggct cgccgggtgc atcacccgga acgtcatcga ccggatcgcc agggagctcg 240

acgccatcag gcgcaacagg atcacctggc tcaagcccta gcgcgtcaac cggcacgggt 300

ccgggtgcct cccctggcac gtccagcacc ggatcacccg gatcgagccc atccgcctca 360

accggaaccg gacccggtac accagggtcg ggaacagcct cctcgtcacc aggctcctca 420

acccccctcgg gagccacggg ttcgcccggt tcgtcaacgc cttccggagc aactggtagc 480

cccggagcat cgccaggaac ttcgagcacg gggtcgcccg cctcgagcga cgcaagattt 540

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tttgtagaat tcacggatca ccttttcaac atcgctaagc caaggccacc ctggatgggt 660

ctgctaggtc ctaccatcca ggctgaggtt tatgatacag tggtcattac acttaagaac 720

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ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct 840

ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

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gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

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agaccatata acatctaccc tcacggaatc actgatgtcc gtcctttgta ttcaaggaga 1920

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gcaaccggtc atcatcacca tcaccattga 5310

<210> 911
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 911
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Gly Ala Pro Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
20 25 30

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
35 40 45

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
50 55 60

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Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
65                  70                  75                  80

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
                85                  90                  95

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
            100                 105                 110

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
        115                 120                 125

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
    130                 135                 140

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
145                 150                 155                 160

Ser Thr Glu Glu Gly Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335
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Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605
```

```
Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880
```

```
Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
        915                 920                 925

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
    930                 935                 940

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
945                 950                 955                 960

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
                965                 970                 975

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly
            980                 985                 990

Thr Ser Ser Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
        995                 1000                1005

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1010                1015                1020

Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser
    1025                1030                1035

Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    1040                1045                1050

Ser Ser  Thr Gly Ser Pro Ala  Ser Ser Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140
```

```
Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395
```

```
Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650
```

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 912
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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tcggaaacgc caggcacaag tgagtctgca actcccgagt ccggacctgg ctccgagcct 180

gccactagcg gctccgagac tccgggaact tccgagagcg ctacaccaga aagcggaccc 240

ggaaccagta ccgaacctag cgagggctct gctccgggca gcccagccgg ctctcctaca 300

tccacggagg agggcacttc cgaatccgcc accccggagt cagggccagg atctgaaccc 360

gctacctcag gcagtgagac gccaggaacg agcgagtccg ctacaccgga gagtgggcca 420

gggagccctg ctggatctcc tacgtccact gaggaagggt caccagcggg ctcgcccacc 480

agcactgaag aaggtgcctc gagcagtgat ctcggtgagc tgcctgtgga cgcaagattt 540

cctcctagag tgccaaaatc ttttccattc aacacctcag tcgtgtacaa aaagactctg 600

tttgtagaat tcacggatca ccttttcaac atcgctaagc caaggccacc ctggatgggt 660

ctgctaggtc ctaccatcca ggctgaggtt tatgatacag tggtcattac acttaagaac 720

atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag 780

ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct 840

ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

ggcctcattg gagccctact agtatgtaga gaagggagtc tggccaagga aaagacacag 1020

accttgcaca aatttatact acttttgct gtatttgatg aagggaaaag ttggcactca 1080

gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg 1200

aaatcagtct attggcatgt gattggaatg ggcaccactc ctgaagtgca ctcaatattc 1260

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ataactttcc ttactgctca aacactcttg atggaccttg gacagtttct actgttttgt 1380

catatctctt cccaccaaca tgatggcatg gaagcttatg tcaaagtaga cagctgtcca 1440

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actgattctg aaatggatgt ggtcaggttt gatgatgaca actctccttc ctttatccaa 1560

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gaggactggg actatgctcc cttagtcctc gcccccgatg acagaagtta taaaagtcaa 1680

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tacacagatg aaacctttaa gactcgtgaa gctattcagc atgaatcagg aatcttggga 1800

cctttacttt atggggaagt tggagacaca ctgttgatta tatttaagaa tcaagcaagc 1860

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gaatttgctc tgtttttcac catctttgat gagaccaaaa gctggtactt cactgaaaat 3900

atggaaagaa actgcagggc tccctgcaat atccagatgg aagatcccac ttttaaagag 3960

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gctcaggatc aaaggattcg atggtatctg ctcagcatgg gcagcaatga aaacatccat 4080

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<210> 913
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 913

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Gly Ala Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
20 25 30

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
35 40 45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
50 55 60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65 70 75 80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
85 90 95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
100 105 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
115 120 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
130 135 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145 150 155 160

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Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430
```

```
Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700
```

```
Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
        915                 920                 925

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
    930                 935                 940

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
945                 950                 955                 960

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
                965                 970                 975
```

```
Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly
            980                 985                 990

Thr Ser Ser Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
        995                 1000                1005

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1010                1015                1020

Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser
    1025                1030                1035

Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    1040                1045                1050

Ser Ser  Thr Gly Ser Pro Ala  Ser Ser Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230
```

```
Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485
```

```
Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1730                1735                1740
```

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 914
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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acggggacgg gaccgggggc gtcaccagga acgtcctcca ccggctcgcc gggtgcatca 180

cccggaacgt catcgaccgg atcgccaggg agctcgacgc catcaggcgc aacaggatca 240

cctggctcaa gccctagcgc gtcaaccggc acgggtccgg gtgcctcccc tggcacgtcc 300

agcaccggat cacccggatc gagcccatcc gcctcaaccg gaaccggacc cggtacacca 360

gggtcgggaa cagcctcctc gtcaccaggc tcctcaaccc cctcgggagc cacgggttcg 420

cccggttcgt caacgccttc cggagcaact ggtagccccg gagcatcgcc aggaacttcg 480

agcacggggt cgcccgcctc gagcagtgat ctcggtgagc tgcctgtgga cgcaagattt 540

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ctgctaggtc ctaccatcca ggctgaggtt tatgatacag tggtcattac acttaagaac 720

atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag 780

ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct 840

ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

ggcctcattg gagccctact agtatgtaga gaagggagtc tggccaagga aaagacacag 1020

accttgcaca aatttatact acttttgct gtatttgatg aagggaaaag ttggcactca 1080

gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg 1200

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ataactttcc ttactgctca aacactcttg atggaccttg gacagtttct actgttttgt 1380

catatctctt cccaccaaca tgatggcatg gaagcttatg tcaaagtaga cagctgtcca 1440

gaggaacccc aactacgaat gaaaaataat gaagaagcgg aagactatga tgatgatctt 1500

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gaggactggg actatgctcc cttagtcctc gcccccgatg acagaagtta taaaagtcaa 1680

tatttgaaca atggccctca gcggattggt aggaagtaca aaaaagtccg atttatggca 1740

tacacagatg aaacctttaa gactcgtgaa gctattcagc atgaatcagg aatcttggga 1800

cctttacttt atggggaagt tggagacaca ctgttgatta tatttaagaa tcaagcaagc 1860

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gatcagaggc aaggagcaga acctagaaaa aactttgtca agcctaatga aaccaaaact 3660

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ggatcacctg gggccgaaac ggccgaacaa aaactcatct cagaagagga tctgtcacct 5280

gcaaccggtc atcatcacca tcaccattga 5310

<210> 915
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 915
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Gly Ala Pro Thr Ser Glu Ser Ala
35 40 45

Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu
50 55 60

Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
65 70 75 80

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala
85 90 95

Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
100 105 110

Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr
115 120 125

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr
130 135 140

Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
145 150 155 160

Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser
165 170 175

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ala Ser Ser Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

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Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525
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Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800
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```
Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
        915                 920                 925

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
    930                 935                 940

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
945                 950                 955                 960

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
                965                 970                 975

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly
            980                 985                 990

Thr Ser Ser Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
        995                 1000                1005

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1010                1015                1020

Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser
    1025                1030                1035

Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    1040                1045                1050

Ser Ser  Thr Gly Ser Pro Ala  Ser Ser Ser Asp Gln  Glu Glu Ile
    1055                1060                1065
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Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320
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Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575
```

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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                 1585                 1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                 1600                 1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                 1615                 1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                 1630                 1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                 1645                 1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                 1660                 1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                 1675                 1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                 1690                 1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                 1705                 1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                 1720                 1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1730                 1735                 1740

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
    1745                 1750                 1755

Ser Pro  Ala Thr Gly His His  His His His His His
    1760                 1765
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<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 916

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ggcgcgccaa catcagagag cgccaccccct gaaagtggtc ccgggagcga gccagccaca      180
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cctacatcca cggaggaggg cacttccgaa tccgccaccc cggagtcagg gccaggatct      420
gaacccgcta cctcaggcag tgagacgcca ggaacgagcg agtccgctac accggagagt      480
gggccaggga gccctgctgg atctcctacg tccactgagg aagggtcacc agcgggctcg      540
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gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa     1140
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<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
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Arg Val Pro Lys Ser Phe Pro Phe Gly Ala Pro Gly Ser Ser Pro Ser
35 40 45

Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
50 55 60

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65 70 75 80

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro

85 90 95

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
100 105 110

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
115 120 125

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser
130 135 140

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145 150 155 160

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
165 170 175

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
260 265 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
275 280 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
290 295 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305 310 315 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
325 330 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
340 345 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln

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        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
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625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
645 650 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
835 840 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
885 890 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln

900 905 910

Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
915 920 925

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
930 935 940

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
945 950 955 960

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
965 970 975

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly
980 985 990

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
995 1000 1005

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
1010 1015 1020

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
1025 1030 1035

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
1040 1045 1050

Ser Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Gln Glu Glu Ile
1055 1060 1065

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
1070 1075 1080

Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln
1085 1090 1095

Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp
1100 1105 1110

Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala
1115 1120 1125

Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu
1130 1135 1140

Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
1145 1150 1155

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val

1160 1165 1170

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1175 1180 1185

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1190 1195 1200

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1205 1210 1215

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1220 1225 1230

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1235 1240 1245

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1250 1255 1260

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1265 1270 1275

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1280 1285 1290

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1295 1300 1305

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1310 1315 1320

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1325 1330 1335

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1340 1345 1350

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1355 1360 1365

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1370 1375 1380

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1385 1390 1395

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1400 1405 1410

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln

1415 1420 1425

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1430 1435 1440

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1445 1450 1455

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1460 1465 1470

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1475 1480 1485

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1490 1495 1500

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1505 1510 1515

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1520 1525 1530

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1535 1540 1545

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1550 1555 1560

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1565 1570 1575

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1580 1585 1590

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1595 1600 1605

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1610 1615 1620

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1625 1630 1635

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn

1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

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<211> 5310
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<213> Artificial Sequence

<220>
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gcatcacccg gaacgtcatc gaccggatcg ccagggagct cgacgccatc aggcgcaaca 300

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Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
35 40 45

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
50 55 60

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
65 70 75 80

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
85 90 95

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
100 105 110

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
115 120 125

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
130 135 140

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
145 150 155 160

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ala Ser Ser
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

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Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
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Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
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Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460
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His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735
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Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
        915                 920                 925

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
    930                 935                 940

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
945                 950                 955                 960

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
                965                 970                 975

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly
            980                 985                 990

Thr Ser Ser Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
        995                 1000                1005
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Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
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Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser
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Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
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Ser Ser  Thr Gly Ser Pro Ala  Ser Ser Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
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Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260
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Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
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Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
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Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
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Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
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Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
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Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
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Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
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Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
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Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
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Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
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Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
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Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
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Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
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Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1730                1735                1740

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
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Ser Pro  Ala Thr Gly His His  His His His His His
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<210> 921
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 921
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285
```

```
Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Gly
385                 390                 395                 400

Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro
                405                 410                 415

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
            420                 425                 430

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
        435                 440                 445

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
    450                 455                 460

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
465                 470                 475                 480

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
                485                 490                 495

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
            500                 505                 510

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
        515                 520                 525

Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    530                 535                 540

Pro Gly Ala Ser Ser Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560
```

```
Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830
```

```
Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
        915                 920                 925

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
    930                 935                 940

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
945                 950                 955                 960

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
                965                 970                 975

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly
            980                 985                 990

Thr Ser Ser Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
        995                 1000                1005

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1010                1015                1020

Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser
    1025                1030                1035

Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    1040                1045                1050

Ser Ser  Thr Gly Ser Pro Ala  Ser Ser Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095
```

```
Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350
```

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Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605
```

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1610 1615 1620

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1625 1630 1635

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

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<211> 5310
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<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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<210> 923
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

```
Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380
```

```
Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
                405                 410                 415

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser
            420                 425                 430

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly
        435                 440                 445

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
    450                 455                 460

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala
465                 470                 475                 480

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly
                485                 490                 495

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
            500                 505                 510

Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala
        515                 520                 525

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
    530                 535                 540

Ser Ser Thr Gly Ser Pro Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655
```

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Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
        915                 920                 925
```

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Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
    930                 935                 940

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
945                 950                 955                 960

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
                965                 970                 975

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly
            980                 985                 990

Thr Ser Ser Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
        995                 1000                1005

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1010                1015                1020

Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser
    1025                1030                1035

Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    1040                1045                1050

Ser Ser  Thr Gly Ser Pro Ala  Ser Ser Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185
```

```
Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                 1195                 1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                 1210                 1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                 1225                 1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                 1240                 1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                 1255                 1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                 1270                 1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                 1285                 1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                 1300                 1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                 1315                 1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                 1330                 1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                 1345                 1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                 1360                 1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                 1375                 1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                 1390                 1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                 1405                 1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                 1420                 1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                 1435                 1440
```

```
Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695
```

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

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<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

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gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

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atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

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<210> 925
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 925

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser

210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
405 410 415

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
420 425 430

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
435 440 445

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
450 455 460

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
465 470 475 480

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser

485 490 495

Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
500 505 510

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
515 520 525

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
530 535 540

Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545 550 555 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
565 570 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
580 585 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
595 600 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
610 615 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
645 650 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn

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        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
        915                 920                 925

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
    930                 935                 940

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
945                 950                 955                 960

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
                965                 970                 975

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly
            980                 985                 990

Thr Ser Ser Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
        995                 1000                1005

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1010                1015                1020

Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser
```

```
    1025                1030                1035

Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    1040                1045                1050

Ser Ser  Thr Gly Ser Pro Ala  Ser Ser Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
```

1280 1285 1290

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1295 1300 1305

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1310 1315 1320

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1325 1330 1335

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1340 1345 1350

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1355 1360 1365

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1370 1375 1380

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1385 1390 1395

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1400 1405 1410

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1415 1420 1425

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1430 1435 1440

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1445 1450 1455

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1460 1465 1470

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1475 1480 1485

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1490 1495 1500

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1505 1510 1515

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1520 1525 1530

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro

1535 1540 1545

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1550 1555 1560

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1565 1570 1575

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1580 1585 1590

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1595 1600 1605

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1610 1615 1620

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1625 1630 1635

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 926
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg gcgcgccaac cagtacggag ccgtccgagg ggagcgcacc aggaagcccg 1260

gctgggagcc cgacttctac cgaagagggt acatctaccg aaccaagtga aggttcagca 1320

ccaggcacct caacagaacc ctctgagggc tcggcgcctg gtacaagtga gtccgccacc 1380

ccagaatccg ggcctgggac aagcacagaa ccttcggaag ggagtgcccc tggaacatcc 1440

gaatcggcaa ccccagaatc agggccagga tctgagcccg cgacttcggg ctccgagacg 1500

cctgggacat ccaccgagcc ctccgaagga tcagccccag gcaccagcac ggagccctct 1560

gagggaagcg cacctggtac cagcgaaagc gcaactcccg aatcaggtcc cggtacgagc 1620

gagtcggcga ccccggagag cgggccaggt gcctcgagcg acagaagtta taaaagtcaa 1680

tatttgaaca atggccctca gcggattggt aggaagtaca aaaaagtccg atttatggca 1740

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cctttacttt atggggaagt tggagacaca ctgttgatta tatttaagaa tcaagcaagc 1860

agaccatata acatctaccc tcacggaatc actgatgtcc gtcctttgta ttcaaggaga 1920

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tataaatgga cagtgactgt agaagatggg ccaactaaat cagatcctcg gtgcctgacc 2040

cgctattact ctagtttcgt taatatggag agagatctag cttcaggact cattggccct 2100

ctcctcatct gctacaaaga atctgtagat caaagaggaa accagataat gtcagacaag 2160

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catttgggac tcctggggcc atatataaga gcagaagttg aagataatat catggtaact 3540

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tgggcttatt tctctgatgt tgacctggaa aaagatgtgc actcaggcct gattggaccc 3780

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tactttacca atatgtttgc cacctggtct ccttcaaaag ctcgacttca cctccaaggg 4860

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caccagattg ccctgaggat ggaggttctg ggctgcgagg cacaggacct ctacggcgcc 5220

ggatcacctg gggccgaaac ggccgaacaa aaactcatct cagaagagga tctgtcacct 5280

gcaaccggtc atcatcacca tcaccattga 5310

<210> 927
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 927
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

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Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320
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Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590
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Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Gly Ala Pro Gly Thr Pro
        755                 760                 765

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser
    770                 775                 780

Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
785                 790                 795                 800

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
                805                 810                 815

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr
            820                 825                 830

Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
        835                 840                 845

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
    850                 855                 860
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Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly
865                 870                 875                 880

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
                885                 890                 895

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Gly Ala Pro Thr Ser Glu Ser Ala Thr Pro Glu
            980                 985                 990

Ser Gly Pro Gly Ser Glu Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly
        995                 1000                1005

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro
    1010                1015                1020

Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr
    1025                1030                1035

Pro Glu  Ser Gly Pro Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    1040                1045                1050

Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    1055                1060                1065

Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala
    1070                1075                1080

Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro
    1085                1090                1095

Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1100                1105                1110

Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    1115                1120                1125
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Ala Ser  Ser Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380
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```
Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                 1390                 1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                 1405                 1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                 1420                 1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                 1435                 1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                 1450                 1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                 1465                 1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                 1480                 1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                 1495                 1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                 1510                 1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                 1525                 1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                 1540                 1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                 1555                 1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                 1570                 1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                 1585                 1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                 1600                 1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                 1615                 1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                 1630                 1635
```

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 928
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

agagatctag cttcaggact cattggccct ctcctcatct gctacaaaga atctgtagat 1680

caaagaggaa accagataat gtcagacaag aggaatgtca tcctgttttc tgtatttgat 1740

gagaaccgaa gctggtacct cacagagaat atacaacgct ttctccccaa tccagctgga 1800

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gtttttgata gtttgcagtt gtcagtttgt ttgcatgagg tggcatactg gtacattcta 1920

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gacagttatg aagatatttc agcatacttg ctgagtaaaa acaatgccat tgaaccaaga 2220

agcttctctc aaaacccacc agtcttgaaa cgccatcaac gggaaataac tcgtactact 2280

cttcagggcg cgccaggtac acccggatcg ggtacagcgt catcgagccc cggtgcgtca 2340

cctggtacgt cgagcacggg gtcgccaggg gcgtcccctg ggacgtcctc aacaggctcg 2400

cccggtgcgt cacccggcac gtcgtccacg ggttcacctg gtagctcccc ttccgcgtcc 2460

actggcaccg ggcctggaac tccggggagc ggcacagcga gctcgtcgcc gggagcatcg 2520

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gccactggat caccagggag ctcgacacct tccggcgcaa cgggatcgcc cggagccagc 2700

ccgggtacgt caagcactgg ctcccctgcc tcgagctcag atcaagagga aatcgattat 2760

gatgatacca tatcagttga aatgaagaag gaagattttg acatttatga tgaggatgaa 2820

aatcagagcc cccgcagctt tcaaaagaaa acacgacact attttattgc tgcagtggag 2880

aggctctggg attatgggat gagtagctcc ccacatgttc taagaaacag ggctcagagt 2940

ggcggcgcgc caacgtccga aagtgctacc cctgagtcag gccctggtag tgagcctgcc 3000

acaagcggaa gcgaaactcc ggggacctca gagtctgcca ctcccgaatc ggggccaggc 3060

tctgaaccgg ccacttcagg gagcgaaaca ccaggaacat cggagagcgc taccccggag 3120

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gccacacccg agtctggccc tggctctcca gcgggctcac ccacgagcac tgaagagggc 3240

tctcccgctg gcagcccaac gtcgacagaa gaaggatcac cagcaggctc ccccacatca 3300

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cccagcgagg ggagtgcacc aggtgcctcg agcagtgtcc ctcagttcaa gaaagttgtt 3420

ttccaggaat ttactgatgg ctcctttact cagcccttat accgtggaga actaaatgaa 3480

catttgggac tcctggggcc atatataaga gcagaagttg aagataatat catggtaact 3540

ttcagaaatc aggcctctcg tccctattcc ttctattcta gccttatttc ttatgaggaa 3600

gatcagaggc aaggagcaga acctagaaaa aactttgtca agcctaatga aaccaaaact 3660

tacttttgga aagtgcaaca tcatatggca cccactaaag atgagtttga ctgcaaagcc 3720

tgggcttatt tctctgatgt tgacctggaa aaagatgtgc actcaggcct gattggaccc 3780

cttctggtct gccacactaa cacactgaac cctgctcatg ggagacaagt gacagtacag 3840

gaatttgctc tgtttttcac catctttgat gagaccaaaa gctggtactt cactgaaaat 3900

atggaaagaa actgcagggc tccctgcaat atccagatgg aagatcccac ttttaaagag 3960

aattatcgct tccatgcaat caatggctac ataatggata cactacctgg cttagtaatg 4020

gctcaggatc aaaggattcg atggtatctg ctcagcatgg gcagcaatga aaacatccat 4080

tctattcatt tcagtggaca tgtgttcact gtacgaaaaa aagaggagta taaaatggca 4140

ctgtacaatc tctatccagg tgtttttgag acagtggaaa tgttaccatc caaagctgga 4200

atttggcggg tggaatgcct tattggcgag catctacatg ctgggatgag cacacttttt 4260

ctggtgtaca gcaataagtg tcagactccc ctgggaatgg cttctggaca cattagagat 4320

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tattccggat caatcaatgc ctggagcacc aaggagccct tttcttggat caaggtggat 4440

ctgttggcac caatgattat tcacggcatc aagacccagg gtgcccgtca gaagttctcc 4500

agcctctaca tctctcagtt tatcatcatg tatagtcttg atgggaagaa gtggcagact 4560

tatcgaggaa attccactgg aaccttaatg gtcttctttg gcaatgtgga ttcatctggg 4620

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cattatagca ttcgcagcac tcttcgcatg gagttgatgg gctgtgattt aaatagttgc 4740

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tactttacca atatgtttgc cacctggtct ccttcaaaag ctcgacttca cctccaaggg 4860

aggagtaatg cctggagacc tcaggtgaat aatccaaaag agtggctgca agtggacttc 4920

cagaagacaa tgaaagtcac aggagtaact actcagggag taaaatctct gcttaccagc 4980

atgtatgtga aggagttcct catctccagc agtcaagatg gccatcagtg gactctcttt 5040

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aactctctag acccaccgtt actgactcgc taccttcgaa ttcaccccca gagttgggtg 5160

caccagattg ccctgaggat ggaggttctg ggctgcgagg cacaggacct ctacggcgcc 5220

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gcaaccggtc atcatcacca tcaccattga 5310

<210> 929
<211> 1772
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 929

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

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Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415
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Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685
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Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Gly Ala Pro Gly Thr Pro
        755                 760                 765

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser
    770                 775                 780

Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
785                 790                 795                 800

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
                805                 810                 815

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr
            820                 825                 830

Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
        835                 840                 845

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
    850                 855                 860

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly
865                 870                 875                 880

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
                885                 890                 895

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960
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Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
            980                 985                 990

Ser Pro Gly Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
        995                 1000                1005

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1010                1015                1020

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1025                1030                1035

Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1040                1045                1050

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1055                1060                1065

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1070                1075                1080

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1085                1090                1095

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1100                1105                1110

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
    1115                1120                1125

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1130                1135                1140

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1145                1150                1155

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1160                1165                1170

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1175                1180                1185

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1190                1195                1200

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1205                1210                1215
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Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1220                1225                1230

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1235                1240                1245

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1250                1255                1260

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1265                1270                1275

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1280                1285                1290

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1295                1300                1305

Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1310                1315                1320

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1325                1330                1335

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1340                1345                1350

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1355                1360                1365

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1370                1375                1380

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1385                1390                1395

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1400                1405                1410

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1415                1420                1425

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1430                1435                1440

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1445                1450                1455

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1460                1465                1470
```

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Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1475                 1480                 1485

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1490                 1495                 1500

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1505                 1510                 1515

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1520                 1525                 1530

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1535                 1540                 1545

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1550                 1555                 1560

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1565                 1570                 1575

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1580                 1585                 1590

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1595                 1600                 1605

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1610                 1615                 1620

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1625                 1630                 1635

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1640                 1645                 1650

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1655                 1660                 1665

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1670                 1675                 1680

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1685                 1690                 1695

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    1700                 1705                 1710

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    1715                 1720                 1725
```

Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala
1730 1735 1740

Gly Ser Pro Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu
1745 1750 1755

Glu Asp Leu Ser Pro Ala Thr Gly His His His His His His
1760 1765 1770

<210> 930
<211> 5319
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 930
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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

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<211> 1769

<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 931

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Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
        35                  40                  45

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
    50                  55                  60

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
65                  70                  75                  80

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
                85                  90                  95

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
            100                 105                 110

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
        115                 120                 125

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
    130                 135                 140

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
145                 150                 155                 160

Ser Thr Glu Glu Gly Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240
```

```
Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510
```

```
Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780
```

```
Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050
```

```
Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
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Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1145                1150                1155

Pro Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    1160                1165                1170

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1175                1180                1185

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1190                1195                1200

Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1205                1210                1215

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1220                1225                1230

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Pro  Ala Gly Ser
    1235                1240                1245

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1250                1255                1260

Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1265                1270                1275

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1280                1285                1290

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ala Ser  Ser Ala Pro
    1295                1300                1305
```

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Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
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Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                 1330                 1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                 1345                 1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                 1360                 1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                 1375                 1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                 1390                 1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                 1405                 1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                 1420                 1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                 1435                 1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                 1450                 1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                 1465                 1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                 1480                 1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                 1495                 1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                 1510                 1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                 1525                 1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                 1540                 1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                 1555                 1560
```

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
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Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
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Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1595 1600 1605

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1610 1615 1620

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
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Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
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Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
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Ser Pro Ala Thr Gly His His His His His His
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<211> 1769
<212> PRT
<213> Artificial Sequence

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35 40 45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
50 55 60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser

65 70 75 80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
85 90 95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
100 105 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
115 120 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
130 135 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145 150 155 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
260 265 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
275 280 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
290 295 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305 310 315 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
325 330 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe

340 345 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
355 360 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
370 375 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385 390 395 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
405 410 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
420 425 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
435 440 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
450 455 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465 470 475 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
485 490 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
500 505 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
515 520 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
530 535 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545 550 555 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
565 570 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
580 585 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
595 600 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn

610 615 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
645 650 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
835 840 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro

885 890 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
900 905 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
915 920 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
930 935 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945 950 955 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
965 970 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
980 985 990

Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
995 1000 1005

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1010 1015 1020

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1025 1030 1035

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1040 1045 1050

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1055 1060 1065

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1070 1075 1080

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1085 1090 1095

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1100 1105 1110

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1115 1120 1125

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1130 1135 1140

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Gly Ala

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Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1175                1180                1185

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1190                1195                1200

Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1205                1210                1215

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1220                1225                1230

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Pro  Ala Gly Ser
    1235                1240                1245

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1250                1255                1260

Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1265                1270                1275

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1280                1285                1290

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ala Ser  Ser Ala Pro
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Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
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Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
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Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
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Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
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Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1415 1420 1425

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1430 1435 1440

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1445 1450 1455

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1460 1465 1470

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1475 1480 1485

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1490 1495 1500

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1505 1510 1515

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1520 1525 1530

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1535 1540 1545

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1550 1555 1560

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1565 1570 1575

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1580 1585 1590

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1595 1600 1605

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1610 1615 1620

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1625 1630 1635

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile

1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
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Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
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<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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20 25 30

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
35 40 45

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
50 55 60

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
65 70 75 80

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
85 90 95

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
100 105 110

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
115 120 125

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
130 135 140

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
145 150 155 160

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ala Ser Ser
165 170 175

```
Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445
```

```
Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720
```

```
Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990
```

```
Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1145                1150                1155

Pro Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    1160                1165                1170

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1175                1180                1185

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1190                1195                1200

Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1205                1210                1215

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1220                1225                1230

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Pro  Ala Gly Ser
    1235                1240                1245
```

```
Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1250                 1255                 1260

Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1265                 1270                 1275

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1280                 1285                 1290

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ala Ser  Ser Ala Pro
    1295                 1300                 1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                 1315                 1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                 1330                 1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                 1345                 1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                 1360                 1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                 1375                 1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                 1390                 1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                 1405                 1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                 1420                 1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                 1435                 1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                 1450                 1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                 1465                 1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                 1480                 1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                 1495                 1500
```

```
Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1730                1735                1740

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
    1745                1750                1755
```

Ser Pro Ala Thr Gly His His His His His His
1760 1765

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<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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gggagcgagc cagccacatc tgggtcggaa acgccaggca caagtgagtc tgcaactccc 180

gagtccggac ctggctccga gcctgccact agcggctccg agactccggg aacttccgag 240

agcgctacac cagaaagcgg acccggaacc agtaccgaac ctagcgaggg ctctgctccg 300

ggcagcccag ccggctctcc tacatccacg gaggagggca cttccgaatc cgccaccccg 360

gagtcagggc caggatctga acccgctacc tcaggcagtg agacgccagg aacgagcgag 420

tccgctacac cggagagtgg gccagggagc cctgctggat ctcctacgtc cactgaggaa 480

gggtcaccag cgggctcgcc caccagcact gaagaaggtg cctcgagcga cgcaagattt 540

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tttgtagaat tcacggatca ccttttcaac atcgctaagc caaggccacc ctggatgggt 660

ctgctaggtc ctaccatcca ggctgaggtt tatgatacag tggtcattac acttaagaac 720

atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag 780

ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct 840

ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

ggcctcattg gagccctact agtatgtaga gaagggagtc tggccaagga aaagacacag 1020

accttgcaca aatttatact acttttgct gtatttgatg aagggaaaag ttggcactca 1080

gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg 1200

aaatcagtct attggcatgt gattggaatg ggcaccactc ctgaagtgca ctcaatattc 1260

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ataactttcc ttactgctca aacactcttg atggaccttg gacagtttct actgttttgt 1380

catatctctt cccaccaaca tgatggcatg gaagcttatg tcaaagtaga cagctgtcca 1440

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attcgctcag ttgccaagaa gcatcctaaa acttgggtac attacattgc tgctgaagag 1620

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tacacagatg aaacctttaa gactcgtgaa gctattcagc atgaatcagg aatcttggga 1800

cctttacttt atggggaagt tggagacaca ctgttgatta tatttaagaa tcaagcaagc 1860

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<210> 937
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 937

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Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Gly Ser Ser
            20                  25                  30

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
        35                  40                  45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
    50                  55                  60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65                  70                  75                  80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
                85                  90                  95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
            100                 105                 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
        115                 120                 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
    130                 135                 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145                 150                 155                 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270
```

```
Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540
```

```
Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815
```

```
Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080
```

```
Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1145                1150                1155

Pro Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    1160                1165                1170

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1175                1180                1185

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1190                1195                1200

Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1205                1210                1215

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1220                1225                1230

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Pro  Ala Gly Ser
    1235                1240                1245

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1250                1255                1260

Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1265                1270                1275

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1280                1285                1290

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ala Ser  Ser Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335
```

```
Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590
```

```
Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1730                1735                1740

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
    1745                1750                1755

Ser Pro  Ala Thr Gly His His  His His His His
    1760                1765
```

<210> 938
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 938

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ccaggttcgt caccctcagc atcaacgggg acgggaccgg gggcgtcacc aggaacgtcc      180

tccaccggct cgccgggtgc atcacccgga acgtcatcga ccggatcgcc agggagctcg      240

acgccatcag gcgcaacagg atcacctggc tcaagcccta gcgcgtcaac cggcacgggt      300
```

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<210> 939
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 939
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Gly Ala Pro Gly Ser Ser Pro Ser
35 40 45

Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
50 55 60

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65 70 75 80

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro
85 90 95

```
Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
            100                 105                 110

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
        115                 120                 125

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser
    130                 135                 140

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145                 150                 155                 160

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
                165                 170                 175

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365
```

```
Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640
```

```
Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910
```

```
Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1145                1150                1155

Pro Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    1160                1165                1170
```

```
Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1175                1180                1185

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1190                1195                1200

Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1205                1210                1215

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1220                1225                1230

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Pro  Ala Gly Ser
    1235                1240                1245

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1250                1255                1260

Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1265                1270                1275

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1280                1285                1290

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ala Ser  Ser Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425
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Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680
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Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 940
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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ggcgcgccag gctccagccc ctccgcgagc acgggaaccg gaccaggttc gtcaccctca 180

gcatcaacgg ggacgggacc gggggcgtca ccaggaacgt cctccaccgg ctcgccgggt 240

gcatcacccg gaacgtcatc gaccggatcg ccagggagct cgacgccatc aggcgcaaca 300

ggatcacctg gctcaagccc tagcgcgtca accggcacgg gtccgggtgc ctcccctggc 360

acgtccagca ccggatcacc cggatcgagc ccatccgcct caaccggaac cggacccggt 420

acaccagggt cgggaacagc ctcctcgtca ccaggctcct caacccccctc gggagccacg 480

ggttcgcccg gttcgtcaac gccttccgga gcaactggta gccccggagc atcgccagga 540

acttcgagca cggggtcgcc cgcctcgagc aacacctcag tcgtgtacaa aaagactctg 600

tttgtagaat tcacggatca ccttttcaac atcgctaagc caaggccacc ctggatgggt 660

ctgctaggtc ctaccatcca ggctgaggtt tatgatacag tggtcattac acttaagaac 720

atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag 780

ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct 840

ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

ggcctcattg gagccctact agtatgtaga gaagggagtc tggccaagga aaagacacag 1020

accttgcaca aatttatact acttttgct gtatttgatg aagggaaaag ttggcactca 1080

gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg 1200

aaatcagtct attggcatgt gattggaatg ggcaccactc ctgaagtgca ctcaatattc 1260

ctcgaaggtc acacatttct tgtgaggaac catcgccagg ctagcttgga aatctcgcca 1320

ataactttcc ttactgctca aacactcttg atggaccttg gacagtttct actgttttgt 1380

catatctctt cccaccaaca tgatggcatg gaagcttatg tcaaagtaga cagctgtcca 1440

gaggaacccc aactacgaat gaaaaataat gaagaagcgg aagactatga tgatgatctt 1500

actgattctg aaatggatgt ggtcaggttt gatgatgaca actctccttc ctttatccaa 1560

attcgctcag ttgccaagaa gcatcctaaa acttgggtac attacattgc tgctgaagag 1620

gaggactggg actatgctcc cttagtcctc gcccccgatg acagaagtta taaaagtcaa 1680

tatttgaaca atggccctca gcggattggt aggaagtaca aaaaagtccg atttatggca 1740

tacacagatg aaacctttaa gactcgtgaa gctattcagc atgaatcagg aatcttggga 1800

cctttacttt atggggaagt tggagacaca ctgttgatta tatttaagaa tcaagcaagc 1860

agaccatata acatctaccc tcacggaatc actgatgtcc gtcctttgta ttcaaggaga 1920

ttaccaaaag gtgtaaaaca tttgaaggat tttccaattc tgccaggaga aatattcaaa 1980

tataaatgga cagtgactgt agaagatggg ccaactaaat cagatcctcg gtgcctgacc 2040

cgctattact ctagtttcgt taatatggag agagatctag cttcaggact cattggccct 2100

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gggtgccaca actcagactt tcggaacaga ggcatgaccg ccttactgaa ggtttctagt 2580

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ctgagtaaaa acaatgccat tgaaccaaga agcttctctc aaaacccacc agtcttgaaa 2700

cgccatcaac gggaaataac tcgtactact cttcagtcag atcaagagga aatcgattat 2760

gatgatacca tatcagttga aatgaagaag gaagattttg acatttatga tgaggatgaa 2820

aatcagagcc cccgcagctt tcaaaagaaa acacgacact attttattgc tgcagtggag 2880

aggctctggg attatgggat gagtagctcc ccacatgttc taagaaacag ggctcagagt 2940

ggcagtgtcc ctcagttcaa gaaagttgtt ttccaggaat ttactgatgg ctcctttact 3000

cagcccttat accgtggaga actaaatgaa catttgggac tcctggggcc atatataaga 3060

gcagaagttg aagataatat catggtaact ttcagaaatc aggcctctcg tccctattcc 3120

ttctattcta gccttatttc ttatgaggaa gatcagaggc aaggagcaga acctagaaaa 3180

aactttgtca agcctaatga aaccaaaact tacttttgga aagtgcaaca tcatatggca 3240

cccactaaag atgagtttga ctgcaaagcc tgggcttatt tctctgatgt tgacctggaa 3300

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cctgctcatg ggagacaagt gacagtacag gaatttgctc tgtttttcac catctttgat 3420

gagaccaaaa gctggtactt cactgaaaat atggaaagaa actgcagggg cgcgccaacg 3480

tccgaaagtg ctacccctga gtcaggccct ggtagtgagc ctgccacaag cggaagcgaa 3540

actccgggga cctcagagtc tgccactccc gaatcggggc caggctctga accggccact 3600

tcagggagcg aaacaccagg aacatcggag agcgctaccc cggagagcgg gccaggaact 3660

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gcaccaggtg cctcgagcgc tccctgcaat atccagatgg aagatcccac ttttaaagag 3960

aattatcgct tccatgcaat caatggctac ataatggata cactacctgg cttagtaatg 4020

gctcaggatc aaaggattcg atggtatctg ctcagcatgg gcagcaatga aaacatccat 4080

tctattcatt tcagtggaca tgtgttcact gtacgaaaaa aagaggagta taaaatggca 4140

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atttggcggg tggaatgcct tattggcgag catctacatg ctgggatgag cacacttttt 4260

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cagaagacaa tgaaagtcac aggagtaact actcagggag taaaatctct gcttaccagc 4980

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aactctctag acccaccgtt actgactcgc taccttcgaa ttcaccccca gagttgggtg 5160

caccagattg ccctgaggat ggaggttctg ggctgcgagg cacaggacct ctacggcgcc 5220

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gcaaccggtc atcatcacca tcaccattga 5310

<210> 941
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 941

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Gly Ala Pro Gly Ser Ser Pro Ser
35 40 45

Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
50 55 60

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65 70 75 80

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro
85 90 95

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
100 105 110

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
115 120 125

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser
130 135 140

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145 150 155 160

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
165 170 175

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu

195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
260 265 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
275 280 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
290 295 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305 310 315 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
325 330 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
340 345 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
355 360 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
370 375 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385 390 395 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
405 410 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
420 425 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
435 440 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
450 455 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro

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465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
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740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
835 840 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
885 890 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
900 905 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
915 920 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
930 935 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945 950 955 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
965 970 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
980 985 990

Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
995 1000 1005

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val

1010 1015 1020

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1025 1030 1035

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1040 1045 1050

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1055 1060 1065

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1070 1075 1080

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1085 1090 1095

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1100 1105 1110

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1115 1120 1125

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1130 1135 1140

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Gly Ala
1145 1150 1155

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala
1160 1165 1170

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
1175 1180 1185

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
1190 1195 1200

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
1205 1210 1215

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala
1220 1225 1230

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
1235 1240 1245

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
1250 1255 1260

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser

1265 1270 1275

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala
1280 1285 1290

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Ala Pro
1295 1300 1305

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1310 1315 1320

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1325 1330 1335

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1340 1345 1350

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1355 1360 1365

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1370 1375 1380

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1385 1390 1395

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1400 1405 1410

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1415 1420 1425

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1430 1435 1440

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1445 1450 1455

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1460 1465 1470

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1475 1480 1485

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1490 1495 1500

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1505 1510 1515

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe

1520 1525 1530

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1535 1540 1545

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1550 1555 1560

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1565 1570 1575

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1580 1585 1590

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1595 1600 1605

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1610 1615 1620

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1625 1630 1635

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 942

<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 942
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ggcgcgccag gctccagccc ctccgcgagc acgggaaccg gaccaggttc gtcaccctca 180

gcatcaacgg ggacgggacc gggggcgtca ccaggaacgt cctccaccgg ctcgccgggt 240

gcatcacccg gaacgtcatc gaccggatcg ccagggagct cgacgccatc aggcgcaaca 300

ggatcacctg gctcaagccc tagcgcgtca accggcacgg gtccgggtgc ctcccctggc 360

acgtccagca ccggatcacc cggatcgagc ccatccgcct caaccggaac cggacccggt 420

acaccagggt cgggaacagc ctcctcgtca ccaggctcct caacccctc gggagccacg 480

ggttcgcccg gttcgtcaac gccttccgga gcaactggta gccccggagc atcgccagga 540

acttcgagca cggggtcgcc cgcctcgagc aacacctcag tcgtgtacaa aaagactctg 600

tttgtagaat tcacggatca ccttttcaac atcgctaagc caaggccacc ctggatgggt 660

ctgctaggtc ctaccatcca ggctgaggtt tatgatacag tggtcattac acttaagaac 720

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<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Gly Ser Ser
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Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
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Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65                  70                  75                  80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
                85                  90                  95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
            100                 105                 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
        115                 120                 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
    130                 135                 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145                 150                 155                 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300
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Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575
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Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845
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Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110
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Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
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Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1145                1150                1155

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala
    1160                1165                1170

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1175                1180                1185

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1190                1195                1200

Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
    1205                1210                1215

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala
    1220                1225                1230

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1235                1240                1245

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1250                1255                1260

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1265                1270                1275

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ala
    1280                1285                1290

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Ala Ser  Ser Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365
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Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620
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Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
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Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
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Ser Pro Ala Thr Gly His His His His His His
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<212> PRT
<213> Artificial Sequence

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<400> 945

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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
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Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
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Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65                  70                  75                  80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
                85                  90                  95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
            100                 105                 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
        115                 120                 125
```

```
Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
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Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145                 150                 155                 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400
```

```
Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670
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```
Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940
```

```
Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
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Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gly Ala Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    1160                1165                1170

Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
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Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1190                1195                1200
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Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala
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Ser Thr  Gly Thr Gly Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
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Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
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Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1250                 1255                 1260

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser
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Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    1280                 1285                 1290

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1295                 1300                 1305

Ala Ser  Ser Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                 1315                 1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                 1330                 1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                 1345                 1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                 1360                 1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                 1375                 1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                 1390                 1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                 1405                 1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                 1420                 1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
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Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                 1450                 1455
```

```
Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
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Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
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Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
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Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
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Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
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Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
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Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
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Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
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Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710
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Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
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Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
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Ser Pro Ala Thr Gly His His His His His His
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cattatagca ttcgcagcac tcttcgcatg gagttgatgg gctgtgattt aaatagttgc 4740
agcatgccat tgggaatgga gagtaaagca atatcagatg cacagattac tgcttcatcc 4800
tactttacca atatgtttgc cacctggtct ccttcaaaag ctcgacttca cctccaaggg 4860
aggagtaatg cctggagacc tcaggtgaat aatccaaaag agtggctgca agtggacttc 4920
cagaagacaa tgaaagtcac aggagtaact actcagggag taaaatctct gcttaccagc 4980
atgtatgtga aggagttcct catctccagc agtcaagatg gccatcagtg gactctcttt 5040
tttcagaatg gcaaagtaaa ggtttttcag ggaaatcaag actccttcac acctgtggtg 5100
aactctctag acccaccgtt actgactcgc taccttcgaa ttcaccccca gagttgggtg 5160
caccagattg ccctgaggat ggaggttctg ggctgcgagg cacaggacct ctacggcgcc 5220
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<210> 947
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 947
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Gly Ala Pro Gly Ser Ser Pro Ser
35 40 45

Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
50 55 60

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65 70 75 80

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro
85 90 95

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
100 105 110

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
115 120 125

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser
130 135 140

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145 150 155 160

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
165 170 175

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

```
Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495
```

```
Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765
```

```
Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035
```

```
Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gly Ala Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    1160                1165                1170

Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1175                1180                1185

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1190                1195                1200

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala
    1205                1210                1215

Ser Thr  Gly Thr Gly Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    1220                1225                1230

Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1235                1240                1245

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1250                1255                1260

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser
    1265                1270                1275

Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    1280                1285                1290
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```
Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1295                1300                1305

Ala Ser  Ser Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545
```

```
Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1730                1735                1740

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
    1745                1750                1755

Ser Pro  Ala Thr Gly His His  His His His His
    1760                1765
```

<210> 948
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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ggcgcgccag gctccagccc ctccgcgagc acgggaaccg gaccaggttc gtcaccctca 180

gcatcaacgg ggacgggacc gggggcgtca ccaggaacgt cctccaccgg ctcgccgggt 240

gcatcacccg gaacgtcatc gaccggatcg ccagggagct cgacgccatc aggcgcaaca 300

ggatcacctg gctcaagccc tagcgcgtca accggcacgg gtccgggtgc ctcccctggc 360

acgtccagca ccggatcacc cggatcgagc ccatccgcct caaccggaac cggacccggt 420

acaccagggt cgggaacagc ctcctcgtca ccaggctcct caacccccctc gggagccacg 480

ggttcgcccg gttcgtcaac gccttccgga gcaactggta gccccggagc atcgccagga 540

acttcgagca cggggtcgcc cgcctcgagc aacacctcag tcgtgtacaa aaagactctg 600

tttgtagaat tcacggatca ccttttcaac atcgctaagc caaggccacc ctggatgggt 660

ctgctaggtc ctaccatcca ggctgaggtt tatgatacag tggtcattac acttaagaac 720

atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag 780

ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct 840

ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

ggcctcattg gagccctact agtatgtaga gaagggagtc tggccaagga aaagacacag 1020

accttgcaca aatttatact acttttgct gtatttgatg aagggaaaag ttggcactca 1080

gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg 1200

aaatcagtct attggcatgt gattggaatg ggcaccactc ctgaagtgca ctcaatattc 1260

ctcgaaggtc acacatttct tgtgaggaac catcgccagg ctagcttgga aatctcgcca 1320

ataactttcc ttactgctca aacactcttg atggaccttg gacagtttct actgttttgt 1380

catatctctt cccaccaaca tgatggcatg gaagcttatg tcaaagtaga cagctgtcca 1440

gaggaacccc aactacgaat gaaaaataat gaagaagcgg aagactatga tgatgatctt 1500

actgattctg aaatggatgt ggtcaggttt gatgatgaca actctccttc ctttatccaa 1560

attcgctcag ttgccaagaa gcatcctaaa acttgggtac attacattgc tgctgaagag 1620

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tatttgaaca atggccctca gcggattggt aggaagtaca aaaaagtccg atttatggca 1740

tacacagatg aaacctttaa gactcgtgaa gctattcagc atgaatcagg aatcttggga 1800

cctttacttt atggggaagt tggagacaca ctgttgatta tatttaagaa tcaagcaagc 1860

agaccatata acatctaccc tcacggaatc actgatgtcc gtcctttgta ttcaaggaga 1920

ttaccaaaag gtgtaaaaca tttgaaggat tttccaattc tgccaggaga aatattcaaa 1980

tataaatgga cagtgactgt agaagatggg ccaactaaat cagatcctcg gtgcctgacc 2040

cgctattact ctagtttcgt taatatggag agagatctag cttcaggact cattggccct 2100

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gtcttcttct ctggatatac cttcaaacac aaaatggtct atgaagacac actcacccta 2460

ttcccattct caggagaaac tgtcttcatg tcgatggaaa acccaggtct atggattctg 2520

gggtgccaca actcagactt tcggaacaga ggcatgaccg ccttactgaa ggtttctagt 2580

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gatgatacca tatcagttga aatgaagaag gaagattttg acatttatga tgaggatgaa 2820

aatcagagcc cccgcagctt tcaaaagaaa acacgacact attttattgc tgcagtggag 2880

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cagcccttat accgtggaga actaaatgaa catttgggac tcctggggcc atatataaga 3060

gcagaagttg aagataatat catggtaact ttcagaaatc aggcctctcg tccctattcc 3120

ttctattcta gccttatttc ttatgaggaa gatcagaggc aaggagcaga acctagaaaa 3180

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gggacatcga gcaccgggtc gccaggagca tcgcccggaa catccagcac aggaagcccc 3780

ggcgcgtcgc ccgggacatc aagcacaggt tccccgggat cgagcacgcc gtccggagcc 3840

actggatcac cagggagctc gacaccttcc ggcgcaacgg gatcgcccgg agccagcccg 3900

ggtacgtcaa gcactggctc ccctgcctcg agccagatgg aagatcccac ttttaaagag 3960

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gctcaggatc aaaggattcg atggtatctg ctcagcatgg gcagcaatga aaacatccat 4080

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atttggcggg tggaatgcct tattggcgag catctacatg ctgggatgag cacacttttt 4260

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ataaaacaca atatttttaa ccctccaatt attgctcgat acatccgttt gcacccaact 4680

cattatagca ttcgcagcac tcttcgcatg gagttgatgg gctgtgattt aaatagttgc 4740

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tactttacca atatgtttgc cacctggtct ccttcaaaag ctcgacttca cctccaaggg 4860

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aactctctag acccaccgtt actgactcgc taccttcgaa ttcaccccca gagttgggtg 5160

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ggatcacctg gggccgaaac ggccgaacaa aaactcatct cagaagagga tctgtcacct 5280

gcaaccggtc atcatcacca tcaccattga 5310

<210> 949
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 949
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Gly Ser Ser
20 25 30

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
35 40 45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser

50 55 60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65 70 75 80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
85 90 95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
100 105 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
115 120 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
130 135 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145 150 155 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
260 265 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
275 280 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
290 295 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305 310 315 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys

325 330 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
340 345 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
355 360 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
370 375 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385 390 395 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
405 410 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
420 425 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
435 440 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
450 455 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465 470 475 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
485 490 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
500 505 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
515 520 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
530 535 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545 550 555 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
565 570 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
580 585 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly

595 600 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
610 615 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
645 650 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
835 840 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu

865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
885 890 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
900 905 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
915 920 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
930 935 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945 950 955 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
965 970 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
980 985 990

Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
995 1000 1005

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1010 1015 1020

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1025 1030 1035

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1040 1045 1050

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1055 1060 1065

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1070 1075 1080

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1085 1090 1095

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1100 1105 1110

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1115 1120 1125

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys

1130 1135 1140

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1145 1150 1155

Cys Asn Ile Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
1160 1165 1170

Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
1175 1180 1185

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
1190 1195 1200

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala
1205 1210 1215

Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
1220 1225 1230

Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
1235 1240 1245

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
1250 1255 1260

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser
1265 1270 1275

Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
1280 1285 1290

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
1295 1300 1305

Ala Ser Ser Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1310 1315 1320

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1325 1330 1335

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1340 1345 1350

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1355 1360 1365

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1370 1375 1380

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys

1385 1390 1395

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1400 1405 1410

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1415 1420 1425

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1430 1435 1440

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1445 1450 1455

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1460 1465 1470

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1475 1480 1485

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1490 1495 1500

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1505 1510 1515

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1520 1525 1530

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1535 1540 1545

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1550 1555 1560

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1565 1570 1575

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1580 1585 1590

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1595 1600 1605

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1610 1615 1620

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1625 1630 1635

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly

1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 950
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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ccaggttcgt caccctcagc atcaacgggg acgggaccgg gggcgtcacc aggaacgtcc 180

tccaccggct cgccgggtgc atcacccgga acgtcatcga ccggatcgcc agggagctcg 240

acgccatcag gcgcaacagg atcacctggc tcaagcccta gcgcgtcaac cggcacgggt 300

ccgggtgcct cccctggcac gtccagcacc ggatcacccg gatcgagccc atccgcctca 360

accggaaccg gacccggtac accagggtcg ggaacagcct cctcgtcacc aggctcctca 420

acccccctcgg gagccacggg ttcgcccggt tcgtcaacgc cttccggagc aactggtagc 480

cccggagcat cgccaggaac ttcgagcacg gggtcgcccg cctcgagcga cgcaagattt 540

cctcctagag tgccaaaatc ttttccattc aacacctcag tcgtgtacaa aaagactctg 600

tttgtagaat tcacggatca ccttttcaac atcgctaagc caaggccacc ctggatgggt 660

ctgctaggtc ctaccatcca ggctgaggtt tatgatacag tggtcattac acttaagaac 720

atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag 780

ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct 840

ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

ggcctcattg gagccctact agtatgtaga gaagggagtc tggccaagga aaagacacag 1020

accttgcaca aatttatact acttttgct gtatttgatg aagggaaaag ttggcactca 1080

gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg 1200

aaatcagtct attggcatgt gattggaatg ggcaccactc ctgaagtgca ctcaatattc 1260

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ataactttcc ttactgctca aacactcttg atggaccttg gacagtttct actgttttgt 1380

catatctctt cccaccaaca tgatggcatg gaagcttatg tcaaagtaga cagctgtcca 1440

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cagaagacaa tgaaagtcac aggagtaact actcagggag taaaatctct gcttaccagc 4980

atgtatgtga aggagttcct catctccagc agtcaagatg gccatcagtg gactctcttt 5040

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aactctctag acccaccgtt actgactcgc taccttcgaa ttcaccccca gagttgggtg 5160

caccagattg ccctgaggat ggaggttctg ggctgcgagg cacaggacct ctacggcgcc 5220

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gcaaccggtc atcatcacca tcaccattga 5310

<210> 951
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 951
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Thr Ser Glu
20 25 30

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
35 40 45

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
50 55 60

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
65 70 75 80

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
85 90 95

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
100 105 110

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
115 120 125

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
130 135 140

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
145 150 155 160

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Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ala Ser Ser
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430
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Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700
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Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975
```

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Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gly Ala Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    1160                1165                1170

Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1175                1180                1185

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1190                1195                1200

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala
    1205                1210                1215

Ser Thr  Gly Thr Gly Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    1220                1225                1230
```

```
Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1235                 1240                 1245

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1250                 1255                 1260

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser
    1265                 1270                 1275

Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    1280                 1285                 1290

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1295                 1300                 1305

Ala Ser  Ser Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                 1315                 1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                 1330                 1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                 1345                 1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                 1360                 1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                 1375                 1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                 1390                 1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                 1405                 1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                 1420                 1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                 1435                 1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                 1450                 1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                 1465                 1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                 1480                 1485
```

```
Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1730                1735                1740
```

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 952
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 952
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ctcggtgagc tgcctgtggg cgcgccaaca tcagagagcg ccacccctga aagtggtccc 120

gggagcgagc cagccacatc tgggtcggaa acgccaggca caagtgagtc tgcaactccc 180

gagtccggac ctggctccga gcctgccact agcggctccg agactccggg aacttccgag 240

agcgctacac cagaaagcgg acccggaacc agtaccgaac ctagcgaggg ctctgctccg 300

ggcagcccag ccggctctcc tacatccacg gaggagggca cttccgaatc cgccaccccg 360

gagtcagggc caggatctga acccgctacc tcaggcagtg agacgccagg aacgagcgag 420

tccgctacac cggagagtgg gccagggagc cctgctggat ctcctacgtc cactgaggaa 480

gggtcaccag cgggctcgcc caccagcact gaagaaggtg cctcgagcga cgcaagattt 540

cctcctagag tgccaaaatc ttttccattc aacacctcag tcgtgtacaa aaagactctg 600

tttgtagaat tcacggatca ccttttcaac atcgctaagc caaggccacc ctggatgggt 660

ctgctaggtc ctaccatcca ggctgaggtt tatgatacag tggtcattac acttaagaac 720

atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag 780

ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct 840

ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

ggcctcattg gagccctact agtatgtaga gaagggagtc tggccaagga aaagacacag 1020

accttgcaca aatttatact acttttgct gtatttgatg aagggaaaag ttggcactca 1080

gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg 1200

aaatcagtct attggcatgt gattggaatg ggcaccactc ctgaagtgca ctcaatattc 1260

ctcgaaggtc acacatttct tgtgaggaac catcgccagg ctagcttgga aatctcgcca 1320

ataactttcc ttactgctca aacactcttg atggaccttg gacagtttct actgttttgt 1380

catatctctt cccaccaaca tgatggcatg gaagcttatg tcaaagtaga cagctgtcca 1440

gaggaacccc aactacgaat gaaaaataat gaagaagcgg aagactatga tgatgatctt 1500

actgattctg aaatggatgt ggtcaggttt gatgatgaca actctccttc ctttatccaa 1560

attcgctcag ttgccaagaa gcatcctaaa acttgggtac attacattgc tgctgaagag 1620

gaggactggg actatgctcc cttagtcctc gcccccgatg acagaagtta taaaagtcaa 1680

tatttgaaca atggccctca gcggattggt aggaagtaca aaaaagtccg atttatggca 1740

tacacagatg aaacctttaa gactcgtgaa gctattcagc atgaatcagg aatcttggga 1800

cctttacttt atggggaagt tggagacaca ctgttgatta tatttaagaa tcaagcaagc 1860

agaccatata acatctaccc tcacggaatc actgatgtcc gtcctttgta ttcaaggaga 1920

ttaccaaaag gtgtaaaaca tttgaaggat tttccaattc tgccaggaga aatattcaaa 1980

tataaatgga cagtgactgt agaagatggg ccaactaaat cagatcctcg gtgcctgacc 2040

cgctattact ctagtttcgt taatatggag agagatctag cttcaggact cattggccct 2100

ctcctcatct gctacaaaga atctgtagat caaagaggaa accagataat gtcagacaag 2160

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tccaacatca tgcacagcat caatggctat gtttttgata gtttgcagtt gtcagtttgt 2340

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ttcccattct caggagaaac tgtcttcatg tcgatggaaa acccaggtct atggattctg 2520

gggtgccaca actcagactt tcggaacaga ggcatgaccg ccttactgaa ggtttctagt 2580

tgtgacaaga acactggtga ttattacgag gacagttatg aagatatttc agcatacttg 2640

ctgagtaaaa acaatgccat tgaaccaaga agcttctctc aaaacccacc agtcttgaaa 2700

cgccatcaac gggaaataac tcgtactact cttcagtcag atcaagagga aatcgattat 2760

gatgatacca tatcagttga aatgaagaag gaagattttg acatttatga tgaggatgaa 2820

aatcagagcc cccgcagctt tcaaaagaaa acacgacact attttattgc tgcagtggag 2880

aggctctggg attatgggat gagtagctcc ccacatgttc taagaaacag ggctcagagt 2940

ggcagtgtcc ctcagttcaa gaaagttgtt ttccaggaat ttactgatgg ctcctttact 3000

cagcccttat accgtggaga actaaatgaa catttgggac tcctggggcc atatataaga 3060

gcagaagttg aagataatat catggtaact ttcagaaatc aggcctctcg tccctattcc 3120

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cccactaaag atgagtttga ctgcaaagcc tgggcttatt tctctgatgt tgacctggaa 3300

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cctgctcatg ggagacaagt gacagtacag gaatttgctc tgtttttcac catctttgat 3420

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gggacatcga gcaccgggtc gccaggagca tcgcccggaa catccagcac aggaagcccc 3780

ggcgcgtcgc ccgggacatc aagcacaggt tccccgggat cgagcacgcc gtccggagcc 3840

actggatcac cagggagctc gacaccttcc ggcgcaacgg gatcgcccgg agccagcccg 3900

ggtacgtcaa gcactggctc ccctgcctcg agccagatgg aagatcccac ttttaaagag 3960

aattatcgct tccatgcaat caatggctac ataatggata cactacctgg cttagtaatg 4020

gctcaggatc aaaggattcg atggtatctg ctcagcatgg gcagcaatga aaacatccat 4080

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atttggcggg tggaatgcct tattggcgag catctacatg ctgggatgag cacacttttt 4260

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ataaaacaca atatttttaa ccctccaatt attgctcgat acatccgttt gcacccaact 4680

cattatagca ttcgcagcac tcttcgcatg gagttgatgg gctgtgattt aaatagttgc 4740

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tactttacca atatgtttgc cacctggtct ccttcaaaag ctcgacttca cctccaaggg 4860

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cagaagacaa tgaaagtcac aggagtaact actcagggag taaaatctct gcttaccagc 4980

atgtatgtga aggagttcct catctccagc agtcaagatg gccatcagtg gactctcttt 5040

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aactctctag acccaccgtt actgactcgc taccttcgaa ttcaccccca gagttgggtg 5160

caccagattg ccctgaggat ggaggttctg ggctgcgagg cacaggacct ctacggcgcc 5220

ggatcacctg gggccgaaac ggccgaacaa aaactcatct cagaagagga tctgtcacct 5280

gcaaccggtc atcatcacca tcaccattga 5310

<210> 953
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 953

```
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1               5                   10                  15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Gly Ala Pro Thr Ser Glu Ser Ala
        35                  40                  45

Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu
    50                  55                  60

Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
65                  70                  75                  80

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala
                85                  90                  95

Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
            100                 105                 110

Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr
        115                 120                 125

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr
    130                 135                 140

Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
145                 150                 155                 160

Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser
                165                 170                 175

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ala Ser Ser Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255
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```
Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525
```

```
Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800
```

```
Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065
```

```
Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gly Ala Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    1160                1165                1170

Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1175                1180                1185

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1190                1195                1200

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala
    1205                1210                1215

Ser Thr  Gly Thr Gly Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    1220                1225                1230

Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1235                1240                1245

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1250                1255                1260

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser
    1265                1270                1275

Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    1280                1285                1290

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1295                1300                1305

Ala Ser  Ser Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320
```

```
Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575
```

```
Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                 1585                 1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                 1600                 1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                 1615                 1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                 1630                 1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                 1645                 1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                 1660                 1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                 1675                 1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                 1690                 1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                 1705                 1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                 1720                 1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1730                 1735                 1740

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
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Ser Pro  Ala Thr Gly His His  His His His His His
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<210> 954
<211> 5310
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<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 954

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ggcgcgccaa catcagagag cgccacccct gaaagtggtc ccgggagcga gccagccaca      180
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ggacccggaa ccagtaccga acctagcgag ggctctgctc cgggcagccc agccggctct 360

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gaacccgcta cctcaggcag tgagacgcca ggaacgagcg agtccgctac accggagagt 480

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ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct 840

ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

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accttgcaca aatttatact acttttgct gtatttgatg aagggaaaag ttggcactca 1080

gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

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catatctctt cccaccaaca tgatggcatg gaagcttatg tcaaagtaga cagctgtcca 1440

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cctttacttt atggggaagt tggagacaca ctgttgatta tatttaagaa tcaagcaagc 1860

agaccatata acatctaccc tcacggaatc actgatgtcc gtcctttgta ttcaaggaga 1920

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cgccatcaac gggaaataac tcgtactact cttcagtcag atcaagagga aatcgattat 2760

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<210> 955
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Gly Ser Ser
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Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
35 40 45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
50 55 60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65 70 75 80

```
Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
                85                  90                  95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
            100                 105                 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
        115                 120                 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
    130                 135                 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145                 150                 155                 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350
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Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620
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Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895
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Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155
```

```
Cys Asn  Ile Gln Met Glu Asp  Pro Gly Ala Pro Gly  Thr Pro Gly
    1160                1165                1170

Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1175                1180                1185

Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1190                1195                1200

Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly
    1205                1210                1215

Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly Pro Gly  Thr Pro Gly
    1220                1225                1230

Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1235                1240                1245

Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1250                1255                1260

Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly
    1265                1270                1275

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr
    1280                1285                1290

Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1295                1300                1305

Ser Thr  Gly Ser Pro Ala Ser  Ser Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410
```

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Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665
```

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

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<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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ccaggttcgt caccctcagc atcaacgggg acgggaccgg gggcgtcacc aggaacgtcc 180

tccaccggct cgccgggtgc atcacccgga acgtcatcga ccggatcgcc agggagctcg 240

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cccggagcat cgccaggaac ttcgagcacg gggtcgcccg cctcgagcga cgcaagattt 540

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ctgctaggtc ctaccatcca ggctgaggtt tatgatacag tggtcattac acttaagaac 720

atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag 780

ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct 840

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gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg 1200

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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
35 40 45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
50 55 60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65 70 75 80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
85 90 95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
100 105 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
115 120 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
130 135 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145 150 155 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr

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Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
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Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
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His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
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Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
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Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Gly Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly
545                 550                 555                 560

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
                565                 570                 575

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
            580                 585                 590

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
        595                 600                 605

Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
    610                 615                 620

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
625                 630                 635                 640

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                645                 650                 655

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            660                 665                 670

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser
        675                 680                 685

Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Leu Ala Pro Asp Asp
    690                 695                 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
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725 730 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
740 745 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
755 760 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
770 775 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785 790 795 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
805 810 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
820 825 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
835 840 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
850 855 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865 870 875 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
885 890 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
900 905 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
915 920 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
930 935 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945 950 955 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
965 970 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
980 985 990

His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val

995 1000 1005

Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr
1010 1015 1020

Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu
1025 1030 1035

Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His Gln
1040 1045 1050

Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
1055 1060 1065

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
1070 1075 1080

Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln
1085 1090 1095

Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp
1100 1105 1110

Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala
1115 1120 1125

Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu
1130 1135 1140

Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
1145 1150 1155

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1160 1165 1170

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1175 1180 1185

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1190 1195 1200

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
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Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1220 1225 1230

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1235 1240 1245

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val

1250 1255 1260

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1265 1270 1275

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1280 1285 1290

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1295 1300 1305

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1310 1315 1320

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1325 1330 1335

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1340 1345 1350

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1355 1360 1365

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1370 1375 1380

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1385 1390 1395

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1400 1405 1410

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1415 1420 1425

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1430 1435 1440

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1445 1450 1455

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1460 1465 1470

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
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Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1490 1495 1500

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp

1505 1510 1515

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1520 1525 1530

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1535 1540 1545

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1550 1555 1560

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1565 1570 1575

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
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Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
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Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
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Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
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1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
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Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
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Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
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Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
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tattccggat caatcaatgc ctggagcacc aaggagccct tttcttggat caaggtggat 4440

ctgttggcac caatgattat tcacggcatc aagacccagg gtgcccgtca gaagttctcc 4500

agcctctaca tctctcagtt tatcatcatg tatagtcttg atgggaagaa gtggcagact 4560

tatcgaggaa attccactgg aaccttaatg gtcttctttg gcaatgtgga ttcatctggg 4620

ataaaacaca atatttttaa ccctccaatt attgctcgat acatccgttt gcacccaact 4680

cattatagca ttcgcagcac tcttcgcatg gagttgatgg gctgtgattt aaatagttgc 4740

agcatgccat tgggaatgga gagtaaagca atatcagatg cacagattac tgcttcatcc 4800

tactttacca atatgtttgc cacctggtct ccttcaaaag ctcgacttca cctccaaggg 4860

aggagtaatg cctggagacc tcaggtgaat aatccaaaag agtggctgca agtggacttc 4920

cagaagacaa tgaaagtcac aggagtaact actcagggag taaaatctct gcttaccagc 4980

atgtatgtga aggagttcct catctccagc agtcaagatg gccatcagtg gactctcttt 5040

tttcagaatg gcaaagtaaa ggtttttcag ggaaatcaag actccttcac acctgtggtg 5100

aactctctag acccaccgtt actgactcgc taccttcgaa ttcaccccca gagttgggtg 5160

caccagattg ccctgaggat ggaggttctg ggctgcgagg cacaggacct ctacggcgcc 5220

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gcaaccggtc atcatcacca tcaccattga 5310

<210> 959
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 959
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Gly Ser Ser
            20                  25                  30

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
        35                  40                  45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
    50                  55                  60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65                  70                  75                  80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
                85                  90                  95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
            100                 105                 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
        115                 120                 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
    130                 135                 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145                 150                 155                 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285
```

```
Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Gly Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly
545                 550                 555                 560
```

```
Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
                565                 570                 575

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
            580                 585                 590

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
        595                 600                 605

Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
    610                 615                 620

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
625                 630                 635                 640

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                645                 650                 655

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            660                 665                 670

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser
        675                 680                 685

Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Leu Ala Pro Asp Asp
    690                 695                 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830
```

```
Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020

Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095
```

```
Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350
```

```
Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                 1360                 1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                 1375                 1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                 1390                 1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                 1405                 1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                 1420                 1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                 1435                 1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                 1450                 1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                 1465                 1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                 1480                 1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                 1495                 1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                 1510                 1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                 1525                 1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                 1540                 1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                 1555                 1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                 1570                 1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                 1585                 1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                 1600                 1605
```

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1610 1615 1620

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1625 1630 1635

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 960
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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ccaggttcgt caccctcagc atcaacgggg acgggaccgg gggcgtcacc aggaacgtcc 180

tccaccggct cgccgggtgc atcacccgga acgtcatcga ccggatcgcc agggagctcg 240

acgccatcag gcgcaacagg atcacctggc tcaagcccta gcgcgtcaac cggcacgggt 300

ccgggtgcct cccctggcac gtccagcacc ggatcacccg gatcgagccc atccgcctca 360

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accggaaccg gacccggtac accagggtcg ggaacagcct cctcgtcacc aggctcctca      420
acccccctcgg gagccacggg ttcgcccggt tcgtcaacgc cttccggagc aactggtagc      480
cccggagcat cgccaggaac ttcgagcacg gggtcgcccg cctcgagcga cgcaagattt      540
cctcctagag tgccaaaatc ttttccattc aacacctcag tcgtgtacaa aaagactctg      600
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ctgctaggtc ctaccatcca ggctgaggtt tatgatacag tggtcattac acttaagaac      720
atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag      780
ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct      840
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ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca      960
ggcctcattg gagccctact agtatgtaga gaagggagtc tggccaagga aaagacacag     1020
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gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa     1140
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ataactttcc ttactgctca aacactcttg atggaccttg gacagtttct actgttttgt     1380
catatctctt cccaccaaca tgatggcatg gaagcttatg tcaaagtaga cagctgtcca     1440
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gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca     2280
ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc     2340
actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat     2400
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gcaaccggtc atcatcacca tcaccattga 5310

<210> 961
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 961

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Gly Ala Pro Gly Ser Ser Pro Ser
35 40 45

Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
50 55 60

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65 70 75 80

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro
85 90 95

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
100 105 110

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Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
        115                 120                 125

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser
    130                 135                 140

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145                 150                 155                 160

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
                165                 170                 175

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380
```

```
Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Gly Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly
545                 550                 555                 560

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
                565                 570                 575

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
            580                 585                 590

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
        595                 600                 605

Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
    610                 615                 620

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
625                 630                 635                 640

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                645                 650                 655
```

```
Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            660                 665                 670

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser
        675                 680                 685

Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Leu Ala Pro Asp Asp
    690                 695                 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925
```

```
Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020

Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185
```

```
Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                 1195                 1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                 1210                 1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                 1225                 1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                 1240                 1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                 1255                 1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                 1270                 1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                 1285                 1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                 1300                 1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                 1315                 1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                 1330                 1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                 1345                 1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                 1360                 1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                 1375                 1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                 1390                 1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                 1405                 1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                 1420                 1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                 1435                 1440
```

```
Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695
```

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 962
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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ggcgcgccag gctccagccc ctccgcgagc acgggaaccg gaccaggttc gtcaccctca 180

gcatcaacgg ggacgggacc gggggcgtca ccaggaacgt cctccaccgg ctcgccgggt 240

gcatcacccg gaacgtcatc gaccggatcg ccagggagct cgacgccatc aggcgcaaca 300

ggatcacctg gctcaagccc tagcgcgtca accggcacgg gtccgggtgc ctcccctggc 360

acgtccagca ccggatcacc cggatcgagc ccatccgcct caaccggaac cggacccggt 420

acaccagggt cgggaacagc ctcctcgtca ccaggctcct caacccctc gggagccacg 480

ggttcgcccg gttcgtcaac gccttccgga gcaactggta gccccggagc atcgccagga 540

acttcgagca cggggtcgcc cgcctcgagc aacacctcag tcgtgtacaa aaagactctg 600

tttgtagaat tcacggatca ccttttcaac atcgctaagc caaggccacc ctggatgggt 660

ctgctaggtc ctaccatcca ggctgaggtt tatgatacag tggtcattac acttaagaac 720

atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag 780

ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct 840

ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

ggcctcattg gagccctact agtatgtaga gaagggagtc tggccaagga aaagacacag 1020

accttgcaca aatttatact acttttgct gtatttgatg aagggaaaag ttggcactca 1080

gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg 1200

aaatcagtct attggcatgt gattggaatg ggcaccactc ctgaagtgca ctcaatattc 1260

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ataactttcc ttactgctca aacactcttg atggaccttg gacagtttct actgttttgt 1380

catatctctt cccaccaaca tgatggcatg gaagcttatg tcaaagtaga cagctgtcca 1440

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attcgctcag ttgccaagaa gcatcctaaa acttgggtac attacattgc tgctgaagag 1620

gaggactggg actatgctcc cttagtcggc gcgccaacca gtacggagcc gtccgagggg 1680

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ccaagtgaag gttcagcacc aggcacctca acagaaccct ctgagggctc ggcgcctggt 1800

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gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 2160

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 2220

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 2280

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 2340

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 2400

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 2460

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 2520

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ggcatgaccg ccttactgaa ggtttctagt tgtgacaaga acactggtga ttattacgag 3060

gacagttatg aagatatttc agcatacttg ctgagtaaaa acaatgccat tgaaccaaga 3120

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gatcagaggc aaggagcaga acctagaaaa aactttgtca agcctaatga aaccaaaact 3660

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gctcaggatc aaaggattcg atggtatctg ctcagcatgg gcagcaatga aaacatccat 4080

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<210> 963
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 963

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Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Gly Ala Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
            20                  25                  30

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
        35                  40                  45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
    50                  55                  60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65                  70                  75                  80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
                85                  90                  95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
            100                 105                 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
        115                 120                 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
    130                 135                 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145                 150                 155                 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205
```

```
Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480
```

```
Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu
545                 550                 555                 560

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
                565                 570                 575

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            580                 585                 590

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
        595                 600                 605

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    610                 615                 620

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
625                 630                 635                 640

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
                645                 650                 655

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
            660                 665                 670

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
        675                 680                 685

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp
    690                 695                 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750
```

```
Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020
```

```
Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275
```

```
Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530
```

```
Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1730                1735                1740

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
    1745                1750                1755

Ser Pro  Ala Thr Gly His His  His His His His His
    1760                1765
```

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<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
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acggggacgg gaccgggggc gtcaccagga acgtcctcca ccggctcgcc gggtgcatca 180

cccggaacgt catcgaccgg atcgccaggg agctcgacgc catcaggcgc aacaggatca 240

cctggctcaa gccctagcgc gtcaaccggc acgggtccgg gtgcctcccc tggcacgtcc 300

agcaccggat cacccggatc gagcccatcc gcctcaaccg gaaccggacc cggtacacca 360

gggtcgggaa cagcctcctc gtcaccaggc tcctcaaccc cctcgggagc cacgggttcg 420

cccggttcgt caacgccttc cggagcaact ggtagccccg gagcatcgcc aggaacttcg 480

agcacggggt cgcccgcctc gagcagtgat ctcggtgagc tgcctgtgga cgcaagattt 540

cctcctagag tgccaaaatc ttttccattc aacacctcag tcgtgtacaa aaagactctg 600

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ctgctaggtc ctaccatcca ggctgaggtt tatgatacag tggtcattac acttaagaac 720

atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag 780

ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct 840

ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

ggcctcattg gagccctact agtatgtaga gaagggagtc tggccaagga aaagacacag 1020

accttgcaca aatttatact acttttgct gtatttgatg aagggaaaag ttggcactca 1080

gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg 1200

aaatcagtct attggcatgt gattggaatg ggcaccactc ctgaagtgca ctcaatattc 1260

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ataactttcc ttactgctca aacactcttg atggaccttg gacagtttct actgttttgt 1380

catatctctt cccaccaaca tgatggcatg gaagcttatg tcaaagtaga cagctgtcca 1440

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ccttcggaag ggagtgcccc tggaacatcc gaatcggcaa ccccagaatc agggccagga 1920

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<210> 965
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 965
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Gly Ala Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
20 25 30

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser

35 40 45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
50 55 60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65 70 75 80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
85 90 95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
100 105 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
115 120 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
130 135 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145 150 155 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
260 265 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
275 280 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
290 295 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser

305 310 315 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
325 330 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
340 345 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
355 360 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
370 375 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385 390 395 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
405 410 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
420 425 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
435 440 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
450 455 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465 470 475 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
485 490 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
500 505 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
515 520 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
530 535 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Gly Ala Ser Pro
545 550 555 560

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
565 570 575

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro

580 585 590

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
595 600 605

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
610 615 620

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
625 630 635 640

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
645 650 655

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
660 665 670

Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
675 680 685

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asp
690 695 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705 710 715 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
725 730 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
740 745 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
755 760 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
770 775 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785 790 795 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
805 810 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
820 825 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
835 840 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn

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    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020

Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
```

1115 1120 1125

Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu
1130 1135 1140

Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
1145 1150 1155

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1160 1165 1170

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1175 1180 1185

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1190 1195 1200

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1205 1210 1215

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1220 1225 1230

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1235 1240 1245

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1250 1255 1260

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1265 1270 1275

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1280 1285 1290

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1295 1300 1305

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1310 1315 1320

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1325 1330 1335

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1340 1345 1350

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1355 1360 1365

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn

1370 1375 1380

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1385 1390 1395

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1400 1405 1410

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1415 1420 1425

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1430 1435 1440

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1445 1450 1455

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1460 1465 1470

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1475 1480 1485

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1490 1495 1500

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1505 1510 1515

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1520 1525 1530

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1535 1540 1545

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1550 1555 1560

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1565 1570 1575

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1580 1585 1590

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1595 1600 1605

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1610 1615 1620

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val

1625 1630 1635

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
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<210> 966
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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acggggacgg gaccgggggc gtcaccagga acgtcctcca ccggctcgcc gggtgcatca 180

cccggaacgt catcgaccgg atcgccaggg agctcgacgc catcaggcgc aacaggatca 240

cctggctcaa gccctagcgc gtcaaccggc acgggtccgg gtgcctcccc tggcacgtcc 300

agcaccggat cacccggatc gagcccatcc gcctcaaccg gaaccggacc cggtacacca 360

gggtcgggaa cagcctcctc gtcaccaggc tcctcaaccc cctcgggagc cacgggttcg 420

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<210> 967
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Gly Ala Pro Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
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Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
35 40 45

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
50 55 60

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
65 70 75 80

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
85 90 95

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
100 105 110

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
115 120 125

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
130 135 140

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Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
145                 150                 155                 160

Ser Thr Glu Glu Gly Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415
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His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Gly Ala Ser Pro
545                 550                 555                 560

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
                565                 570                 575

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            580                 585                 590

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
        595                 600                 605

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
    610                 615                 620

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
625                 630                 635                 640

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
                645                 650                 655

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
            660                 665                 670

Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
        675                 680                 685
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Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asp
    690                 695                 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960
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Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
               965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
           980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
       995                1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
   1010                1015                1020

Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
   1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Pro Pro Val Leu Lys  Arg His Gln
   1040                1045                1050

Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
   1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
   1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
   1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
   1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
   1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
   1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
   1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
   1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
   1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
   1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
   1205                1210                1215
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Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                 1225                 1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                 1240                 1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                 1255                 1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                 1270                 1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                 1285                 1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                 1300                 1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                 1315                 1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                 1330                 1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                 1345                 1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                 1360                 1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                 1375                 1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                 1390                 1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                 1405                 1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                 1420                 1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                 1435                 1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                 1450                 1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                 1465                 1470
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Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725
```

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 968
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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tcggaaacgc caggcacaag tgagtctgca actcccgagt ccggacctgg ctccgagcct 180

gccactagcg gctccgagac tccgggaact tccgagagcg ctacaccaga aagcggaccc 240

ggaaccagta ccgaacctag cgagggctct gctccgggca gcccagccgg ctctcctaca 300

tccacggagg agggcacttc cgaatccgcc accccggagt cagggccagg atctgaaccc 360

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gggagccctg ctggatctcc tacgtccact gaggaagggt caccagcggg ctcgcccacc 480

agcactgaag aaggtgcctc gagcagtgat ctcggtgagc tgcctgtgga cgcaagattt 540

cctcctagag tgccaaaatc ttttccattc aacacctcag tcgtgtacaa aaagactctg 600

tttgtagaat tcacggatca ccttttcaac atcgctaagc caaggccacc ctggatgggt 660

ctgctaggtc ctaccatcca ggctgaggtt tatgatacag tggtcattac acttaagaac 720

atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag 780

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accttgcaca aatttatact acttttgct gtatttgatg aagggaaaag ttggcactca 1080

gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg 1200

aaatcagtct attggcatgt gattggaatg ggcaccactc ctgaagtgca ctcaatattc 1260

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<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 969

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240
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Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510
```

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Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Gly Ala Pro Gly Thr Pro
        755                 760                 765

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser
    770                 775                 780
```

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Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
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Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
                805                 810                 815

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr
            820                 825                 830

Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
        835                 840                 845

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
    850                 855                 860

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly
865                 870                 875                 880

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
                885                 890                 895

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050
```

```
Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
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Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1145                1150                1155

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala
    1160                1165                1170

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1175                1180                1185

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1190                1195                1200

Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
    1205                1210                1215

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala
    1220                1225                1230

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1235                1240                1245

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1250                1255                1260

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1265                1270                1275

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ala
    1280                1285                1290

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Ala Ser  Ser Ala Pro
    1295                1300                1305
```

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Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
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Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                 1330                 1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                 1345                 1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                 1360                 1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                 1375                 1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                 1390                 1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                 1405                 1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                 1420                 1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                 1435                 1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                 1450                 1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                 1465                 1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                 1480                 1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                 1495                 1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                 1510                 1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                 1525                 1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                 1540                 1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                 1555                 1560
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Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
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Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
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Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
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Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
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Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
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Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
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Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
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Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
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Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
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Ser Pro Ala Thr Gly His His His His His His
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gagaccaaaa gctggtactt cactgaaaat atggaaagaa actgcagggg cgcgccaggt 3480

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<210> 971
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 971
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

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Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335
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Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605
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```
Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Gly Ala Pro Gly Thr Pro
        755                 760                 765

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser
    770                 775                 780

Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
785                 790                 795                 800

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
                805                 810                 815

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr
            820                 825                 830

Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
        835                 840                 845

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
    850                 855                 860

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly
865                 870                 875                 880
```

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Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
                885                 890                 895

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140
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Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1145                1150                1155

Pro Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    1160                1165                1170

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1175                1180                1185

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1190                1195                1200

Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1205                1210                1215

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1220                1225                1230

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Pro  Ala Gly Ser
    1235                1240                1245

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1250                1255                1260

Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1265                1270                1275

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1280                1285                1290

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ala Ser  Ser Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395
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Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650
```

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 972
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

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gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

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catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

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gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

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cccggtgcgt cacccggcac gtcgtccacg ggttcacctg gtagctcccc ttccgcgtcc 2460

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<210> 973
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 973

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu

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                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
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435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu

705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
740 745 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Gly Ala Pro Gly Thr Pro
755 760 765

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser
770 775 780

Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
785 790 795 800

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
805 810 815

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr
820 825 830

Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
835 840 845

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
850 855 860

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly
865 870 875 880

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
885 890 895

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
900 905 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
915 920 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
930 935 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945 950 955 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
965 970 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln

980 985 990

Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
995 1000 1005

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1010 1015 1020

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1025 1030 1035

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1040 1045 1050

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1055 1060 1065

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1070 1075 1080

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1085 1090 1095

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1100 1105 1110

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1115 1120 1125

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1130 1135 1140

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1145 1150 1155

Cys Asn Ile Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
1160 1165 1170

Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
1175 1180 1185

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
1190 1195 1200

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala
1205 1210 1215

Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
1220 1225 1230

Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro

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    1235                1240                1245

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1250                1255                1260

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser
    1265                1270                1275

Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    1280                1285                1290

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1295                1300                1305

Ala Ser  Ser Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
```

1490 1495 1500

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1505 1510 1515

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1520 1525 1530

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1535 1540 1545

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1550 1555 1560

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1565 1570 1575

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1580 1585 1590

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1595 1600 1605

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1610 1615 1620

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1625 1630 1635

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu

1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

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<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

agagatctag cttcaggact cattggccct ctcctcatct gctacaaaga atctgtagat 1680

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cttcagggcg cgccaggtac acccggatcg ggtacagcgt catcgagccc cggtgcgtca 2340

cctggtacgt cgagcacggg gtcgccaggg gcgtcccctg ggacgtcctc aacaggctcg 2400

cccggtgcgt cacccggcac gtcgtccacg ggttcacctg gtagctcccc ttccgcgtcc 2460

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gccactggat caccagggag ctcgacacct tccggcgcaa cgggatcgcc cggagccagc 2700

ccgggtacgt caagcactgg ctcccctgcc tcgagctcag atcaagagga aatcgattat 2760

gatgatacca tatcagttga aatgaagaag gaagattttg acatttatga tgaggatgaa 2820

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aggctctggg attatgggat gagtagctcc ccacatgttc taagaaacag ggctcagagt 2940

ggcagtgtcc ctcagttcaa gaaagttgtt ttccaggaat ttactgatgg ctcctttact 3000

cagcccttat accgtggaga actaaatgaa catttgggac tcctggggcc atatataaga 3060

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aattatcgct tccatgcaat caatggctac ataatggata cactacctgg cttagtaatg 4020

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atttggcggg tggaatgcct tattggcgag catctacatg ctgggatgag cacacttttt 4260

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tactttacca atatgtttgc cacctggtct ccttcaaaag ctcgacttca cctccaaggg 4860

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cagaagacaa tgaaagtcac aggagtaact actcagggag taaaatctct gcttaccagc 4980

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aactctctag acccaccgtt actgactcgc taccttcgaa ttcaccccca gagttgggtg 5160

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gcaaccggtc atcatcacca tcaccattga 5310

<210> 975
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 975
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Gly Ala Pro Gly Ser Ser Pro Ser
35 40 45

Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
50 55 60

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65 70 75 80

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro
85 90 95

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
100 105 110

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
115 120 125

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser
130 135 140

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145 150 155 160

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
165 170 175

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
260 265 270

```
Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540
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Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815
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Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Gly Ala Pro Thr Ser Glu Ser Ala Thr
                965                 970                 975

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
            980                 985                 990

Pro Gly Thr Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
        995                 1000                1005

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1010                1015                1020

Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1040                1045                1050

Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr Glu Glu  Gly Ser Pro
    1055                1060                1065

Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Ser Pro  Ala Gly Ser
    1070                1075                1080
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Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1085                 1090                 1095

Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1100                 1105                 1110

Gly Ala  Ser Ser Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                 1120                 1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                 1135                 1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                 1150                 1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                 1165                 1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                 1180                 1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                 1195                 1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                 1210                 1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                 1225                 1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                 1240                 1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                 1255                 1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                 1270                 1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                 1285                 1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                 1300                 1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                 1315                 1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                 1330                 1335
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Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590
```

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1595 1600 1605

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1610 1615 1620

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1625 1630 1635

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 976
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 976
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ctcggtgagc tgcctgtgga cgcaagattt cctcctagag tgccaaaatc ttttccattc 120

ggcgcgccag gctccagccc ctccgcgagc acgggaaccg gaccaggttc gtcaccctca 180

gcatcaacgg ggacgggacc gggggcgtca ccaggaacgt cctccaccgg ctcgccgggt 240

gcatcacccg gaacgtcatc gaccggatcg ccagggagct cgacgccatc aggcgcaaca 300

ggatcacctg gctcaagccc tagcgcgtca accggcacgg gtccgggtgc ctcccctggc 360

acgtccagca ccggatcacc cggatcgagc ccatccgcct caaccggaac cggacccggt 420

acaccagggt cgggaacagc ctcctcgtca ccaggctcct caacccccctc gggagccacg 480

ggttcgcccg gttcgtcaac gccttccgga gcaactggta gccccggagc atcgccagga 540

acttcgagca cggggtcgcc cgcctcgagc aacacctcag tcgtgtacaa aaagactctg 600

tttgtagaat tcacggatca ccttttcaac atcgctaagc caaggccacc ctggatgggt 660

ctgctaggtc ctaccatcca ggctgaggtt tatgatacag tggtcattac acttaagaac 720

atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag 780

ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct 840

ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

ggcctcattg gagccctact agtatgtaga gaagggagtc tggccaagga aaagacacag 1020

accttgcaca aatttatact acttttgct gtatttgatg aagggaaaag ttggcactca 1080

gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg 1200

aaatcagtct attggcatgt gattggaatg ggcaccactc ctgaagtgca ctcaatattc 1260

ctcgaaggtc acacatttct tgtgaggaac catcgccagg ctagcttgga aatctcgcca 1320

ataactttcc ttactgctca aacactcttg atggaccttg gacagtttct actgttttgt 1380

catatctctt cccaccaaca tgatggcatg gaagcttatg tcaaagtaga cagctgtcca 1440

gaggaacccc aactacgaat gaaaaataat gaagaagcgg aagactatga tgatgatctt 1500

actgattctg aaatggatgt ggtcaggttt gatgatgaca actctccttc ctttatccaa 1560

attcgctcag ttgccaagaa gcatcctaaa acttgggtac attacattgc tgctgaagag 1620

gaggactggg actatgctcc cttagtcctc gcccccgatg acagaagtta taaaagtcaa 1680

tatttgaaca atggccctca gcggattggt aggaagtaca aaaaagtccg atttatggca 1740

tacacagatg aaacctttaa gactcgtgaa gctattcagc atgaatcagg aatcttggga 1800

cctttacttt atggggaagt tggagacaca ctgttgatta tatttaagaa tcaagcaagc 1860

agaccatata acatctaccc tcacggaatc actgatgtcc gtcctttgta ttcaaggaga 1920

ttaccaaaag gtgtaaaaca tttgaaggat tttccaattc tgccaggaga aatattcaaa 1980

tataaatgga cagtgactgt agaagatggg ccaactaaat cagatcctcg gtgcctgacc 2040

cgctattact ctagtttcgt taatatggag agagatctag cttcaggact cattggccct 2100

ctcctcatct gctacaaaga atctgtagat caaagaggaa accagataat gtcagacaag 2160

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gggtgccaca actcagactt tcggaacaga ggcatgaccg ccttactgaa ggtttctagt 2580

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cgccatcaac gggaaataac tcgtactact cttcagtcag atcaagagga aatcgattat 2760

gatgatacca tatcagttga aatgaagaag gaagattttg acatttatga tgaggatgaa 2820

aatcagagcc cccgcagctt tcaaaagaaa acacgacact attttattgc tgcagtggag 2880

aggctctggg attatgggat gggcgcgcca acgtccgaaa gtgctacccc tgagtcaggc 2940

cctggtagtg agcctgccac aagcggaagc gaaactccgg ggacctcaga gtctgccact 3000

cccgaatcgg ggccaggctc tgaaccggcc acttcaggga gcgaaacacc aggaacatcg 3060

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cctggtacaa gcgagtccgc cacacccgag tctggccctg gctctccagc gggctcaccc 3180

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gcaggctccc ccacatcaac agaggagggt acatcagaat ctgctactcc cgagagtgga 3300

cccggtacct ccactgagcc cagcgagggg agtgcaccag gtgcctcgag cagtagctcc 3360

ccacatgttc taagaaacag ggctcagagt ggcagtgtcc ctcagttcaa gaaagttgtt 3420

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catttgggac tcctggggcc atatataaga gcagaagttg aagataatat catggtaact 3540

ttcagaaatc aggcctctcg tccctattcc ttctattcta gccttatttc ttatgaggaa 3600

gatcagaggc aaggagcaga acctagaaaa aactttgtca agcctaatga aaccaaaact 3660

tacttttgga aagtgcaaca tcatatggca cccactaaag atgagtttga ctgcaaagcc 3720

tgggcttatt tctctgatgt tgacctggaa aaagatgtgc actcaggcct gattggaccc 3780

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gaatttgctc tgtttttcac catctttgat gagaccaaaa gctggtactt cactgaaaat 3900

atggaaagaa actgcagggc tccctgcaat atccagatgg aagatcccac ttttaaagag 3960

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gctcaggatc aaaggattcg atggtatctg ctcagcatgg gcagcaatga aaacatccat 4080

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atttggcggg tggaatgcct tattggcgag catctacatg ctgggatgag cacacttttt 4260

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tatcgaggaa attccactgg aaccttaatg gtcttctttg gcaatgtgga ttcatctggg 4620

ataaaacaca atatttttaa ccctccaatt attgctcgat acatccgttt gcacccaact 4680

cattatagca ttcgcagcac tcttcgcatg gagttgatgg gctgtgattt aaatagttgc 4740

agcatgccat tgggaatgga gagtaaagca atatcagatg cacagattac tgcttcatcc 4800

tactttacca atatgtttgc cacctggtct ccttcaaaag ctcgacttca cctccaaggg 4860

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cagaagacaa tgaaagtcac aggagtaact actcagggag taaaatctct gcttaccagc 4980

atgtatgtga aggagttcct catctccagc agtcaagatg gccatcagtg gactctcttt 5040

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aactctctag acccaccgtt actgactcgc taccttcgaa ttcaccccca gagttgggtg 5160

caccagattg ccctgaggat ggaggttctg ggctgcgagg cacaggacct ctacggcgcc 5220

ggatcacctg gggccgaaac ggccgaacaa aaactcatct cagaagagga tctgtcacct 5280

gcaaccggtc atcatcacca tcaccattga 5310

<210> 977
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 977
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Thr Ser Glu
20 25 30

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
35 40 45

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
50 55 60

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
65 70 75 80

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
85 90 95

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Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
            100                 105                 110

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
        115                 120                 125

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
    130                 135                 140

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
145                 150                 155                 160

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ala Ser Ser
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365
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Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640
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Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910
```

```
Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Gly Ala Pro Thr Ser Glu Ser Ala Thr
                965                 970                 975

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
            980                 985                 990

Pro Gly Thr Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
        995                 1000                1005

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1010                1015                1020

Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1040                1045                1050

Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr Glu Glu  Gly Ser Pro
    1055                1060                1065

Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Ser Pro  Ala Gly Ser
    1070                1075                1080

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1085                1090                1095

Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1100                1105                1110

Gly Ala  Ser Ser Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170
```

```
Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425
```

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Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
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Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680
```

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Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                 1690                 1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                 1705                 1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                 1720                 1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1730                 1735                 1740

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
    1745                 1750                 1755

Ser Pro  Ala Thr Gly His His  His His His His
    1760                 1765
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<210> 978
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 978

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ggcagcccag ccggctctcc tacatccacg gaggagggca cttccgaatc cgccaccccg      360

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<210> 979
<211> 1772
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Gly Ser Ser
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Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
35 40 45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
50 55 60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65 70 75 80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
85 90 95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
100 105 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
115 120 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
130 135 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145 150 155 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

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Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460
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His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735
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Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
            980                 985                 990

Ser Pro Gly Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
        995                 1000                1005
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Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
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Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1025                 1030                 1035

Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1040                 1045                 1050

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1055                 1060                 1065

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1070                 1075                 1080

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1085                 1090                 1095

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1100                 1105                 1110

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
    1115                 1120                 1125

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1130                 1135                 1140

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1145                 1150                 1155

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1160                 1165                 1170

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1175                 1180                 1185

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1190                 1195                 1200

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1205                 1210                 1215

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1220                 1225                 1230

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1235                 1240                 1245

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1250                 1255                 1260
```

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Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
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Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1280                1285                1290

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1295                1300                1305

Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1310                1315                1320

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1325                1330                1335

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1340                1345                1350

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1355                1360                1365

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1370                1375                1380

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1385                1390                1395

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1400                1405                1410

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1415                1420                1425

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1430                1435                1440

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1445                1450                1455

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1460                1465                1470

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1475                1480                1485

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1490                1495                1500

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1505                1510                1515
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Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1520                1525                1530

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1535                1540                1545

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1550                1555                1560

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1565                1570                1575

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1580                1585                1590

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1595                1600                1605

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1610                1615                1620

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1625                1630                1635

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1640                1645                1650

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1655                1660                1665

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1670                1675                1680

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1685                1690                1695

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    1700                1705                1710

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    1715                1720                1725

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    1730                1735                1740

Gly Ser  Pro Gly Ala Glu Thr  Ala Glu Gln Lys Leu  Ile Ser Glu
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Glu Asp  Leu Ser Pro Ala Thr  Gly His His His His  His His
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<211> 5319
<212> DNA
<213> Artificial Sequence

<220>
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ccaggttcgt caccctcagc atcaacgggg acgggaccgg gggcgtcacc aggaacgtcc 180

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Arg Val Pro Lys Ser Phe Pro Phe Gly Ala Pro Gly Ser Ser Pro Ser
35 40 45

Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
50 55 60

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65 70 75 80

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro
85 90 95

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
100 105 110

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
115 120 125

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser
130 135 140

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145 150 155 160

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
165 170 175

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
260 265 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
275 280 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu

290 295 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305 310 315 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
325 330 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
340 345 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
355 360 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
370 375 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385 390 395 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
405 410 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
420 425 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
435 440 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
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His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465 470 475 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
485 490 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
500 505 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
515 520 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
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Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545 550 555 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val

565 570 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
580 585 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
595 600 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
610 615 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
645 650 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg

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Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
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Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
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Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
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Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
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Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
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Arg Ala Gln Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
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Ser Pro Gly Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
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Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
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Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
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Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
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Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1085                1090                1095

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
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Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser
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Ser Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val
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1145 1150 1155

Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
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Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu
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Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser
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Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
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Asn Tyr Arg Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu
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Pro Gly Leu Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu
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Leu Ser Met Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser

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Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
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Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
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His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
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Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
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Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
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Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
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Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
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Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
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Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
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Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
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Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
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Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
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His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
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Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
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Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
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Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
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Thr Gln Gly Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu
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Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg
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Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu
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Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala
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<211> 1772
<212> PRT
<213> Artificial Sequence

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Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
35 40 45

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
50 55 60

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
65 70 75 80

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
85 90 95

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
100 105 110

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
115 120 125

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Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
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Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
145                 150                 155                 160

Ser Thr Glu Glu Gly Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400
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Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670
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Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940
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Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
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Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
            980                 985                 990

Ser Pro Gly Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
        995                 1000                1005

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1010                1015                1020

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
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Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1040                1045                1050

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1055                1060                1065

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1070                1075                1080

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1085                1090                1095

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1100                1105                1110

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
    1115                1120                1125

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1130                1135                1140

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1145                1150                1155

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1160                1165                1170

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1175                1180                1185

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1190                1195                1200
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Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
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Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1220                 1225                 1230

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1235                 1240                 1245

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1250                 1255                 1260

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1265                 1270                 1275

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1280                 1285                 1290

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1295                 1300                 1305

Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1310                 1315                 1320

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1325                 1330                 1335

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1340                 1345                 1350

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1355                 1360                 1365

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
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Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1385                 1390                 1395

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1400                 1405                 1410

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1415                 1420                 1425

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1430                 1435                 1440

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1445                 1450                 1455
```

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Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
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Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1475                 1480                 1485

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1490                 1495                 1500

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1505                 1510                 1515

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1520                 1525                 1530

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1535                 1540                 1545

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1550                 1555                 1560

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1565                 1570                 1575

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1580                 1585                 1590

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1595                 1600                 1605

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1610                 1615                 1620

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
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Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1640                 1645                 1650

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1655                 1660                 1665

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1670                 1675                 1680

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1685                 1690                 1695

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    1700                 1705                 1710
```

Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu
1715 1720 1725

Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala
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Gly Ser Pro Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu
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Glu Asp Leu Ser Pro Ala Thr Gly His His His His His His
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aatagttgca gcatgccatt gggaatggag agtaaagcaa tatcagatgc acagattact 4800

gcttcatcct actttaccaa tatgtttgcc acctggtctc cttcaaaagc tcgacttcac 4860

ctccaaggga ggagtaatgc ctggagacct caggtgaata atccaaaaga gtggctgcaa 4920

gtggacttcc agaagacaat gaaagtcaca ggagtaacta ctcagggagt aaaatctctg 4980

cttaccagca tgtatgtgaa ggagttcctc atctccagca gtcaagatgg ccatcagtgg 5040

actctctttt ttcagaatgg caaagtaaag gtttttcagg gaaatcaaga ctccttcaca 5100

cctgtggtga actctctaga cccaccgtta ctgactcgct accttcgaat tcacccccag 5160

agttgggtgc accagattgc cctgaggatg gaggttctgg gctgcgaggc acaggacctc 5220

tacggcgccg gatcacctgg ggccgaaacg gccgaacaaa aactcatctc agaagaggat 5280

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<210> 985
<211> 1772
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 985
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Thr Ser Glu
20 25 30

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
35 40 45

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
50 55 60

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
65 70 75 80

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
85 90 95

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
100 105 110

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
115 120 125

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
130 135 140

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
145 150 155 160

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ala Ser Ser
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

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Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495
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Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765
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Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
            980                 985                 990

Ser Pro Gly Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
        995                 1000                1005

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1010                1015                1020

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1025                1030                1035
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Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1040                1045                1050

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1055                1060                1065

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1070                1075                1080

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1085                1090                1095

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1100                1105                1110

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
    1115                1120                1125

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1130                1135                1140

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1145                1150                1155

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1160                1165                1170

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1175                1180                1185

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1190                1195                1200

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1205                1210                1215

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1220                1225                1230

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1235                1240                1245

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1250                1255                1260

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1265                1270                1275

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1280                1285                1290
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Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1295                1300                1305

Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1310                1315                1320

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1325                1330                1335

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1340                1345                1350

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1355                1360                1365

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1370                1375                1380

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1385                1390                1395

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1400                1405                1410

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1415                1420                1425

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1430                1435                1440

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1445                1450                1455

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1460                1465                1470

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1475                1480                1485

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1490                1495                1500

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1505                1510                1515

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1520                1525                1530

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1535                1540                1545
```

Phe Asn Pro Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr
1550 1555 1560

His Tyr Ser Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys
1565 1570 1575

Asp Leu Asn Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala
1580 1585 1590

Ile Ser Asp Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met
1595 1600 1605

Phe Ala Thr Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly
1610 1615 1620

Arg Ser Asn Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp
1625 1630 1635

Leu Gln Val Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr
1640 1645 1650

Thr Gln Gly Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu
1655 1660 1665

Phe Leu Ile Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe
1670 1675 1680

Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser
1685 1690 1695

Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg
1700 1705 1710

Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu
1715 1720 1725

Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala
1730 1735 1740

Gly Ser Pro Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu
1745 1750 1755

Glu Asp Leu Ser Pro Ala Thr Gly His His His His His His
1760 1765 1770

<210> 986
<211> 5319
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic

polynucleotide

<400> 986

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gggagcgagc cagccacatc tgggtcggaa acgccaggca caagtgagtc tgcaactccc 180

gagtccggac ctggctccga gcctgccact agcggctccg agactccggg aacttccgag 240

agcgctacac cagaaagcgg acccggaacc agtaccgaac ctagcgaggg ctctgctccg 300

ggcagcccag ccggctctcc tacatccacg gaggagggca cttccgaatc cgccaccccg 360

gagtcagggc caggatctga acccgctacc tcaggcagtg agacgccagg aacgagcgag 420

tccgctacac cggagagtgg gccagggagc cctgctggat ctcctacgtc cactgaggaa 480

gggtcaccag cgggctcgcc caccagcact gaagaaggtg cctcgagcga cgcaagattt 540

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atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag 780

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ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

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<210> 987
<211> 1772
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 987
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Gly Ala Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
20 25 30

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
35 40 45

```
Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
    50                  55                  60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65                  70                  75                  80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
                85                  90                  95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
            100                 105                 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
        115                 120                 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
    130                 135                 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145                 150                 155                 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320
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Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590
```

```
Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860
```

```
Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
            980                 985                 990

Ser Pro Gly Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
        995                 1000                1005

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1010                1015                1020

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1025                1030                1035

Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1040                1045                1050

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1055                1060                1065

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1070                1075                1080

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1085                1090                1095

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1100                1105                1110

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
    1115                1120                1125
```

```
Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1130                1135                1140

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1145                1150                1155

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1160                1165                1170

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1175                1180                1185

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1190                1195                1200

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1205                1210                1215

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1220                1225                1230

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1235                1240                1245

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1250                1255                1260

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1265                1270                1275

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1280                1285                1290

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1295                1300                1305

Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1310                1315                1320

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1325                1330                1335

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1340                1345                1350

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1355                1360                1365

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1370                1375                1380
```

```
Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1385                1390                1395

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1400                1405                1410

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1415                1420                1425

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1430                1435                1440

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1445                1450                1455

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1460                1465                1470

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1475                1480                1485

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1490                1495                1500

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1505                1510                1515

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1520                1525                1530

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1535                1540                1545

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1550                1555                1560

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1565                1570                1575

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1580                1585                1590

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1595                1600                1605

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1610                1615                1620

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1625                1630                1635
```

Leu Gln Val Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr
1640 1645 1650

Thr Gln Gly Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu
1655 1660 1665

Phe Leu Ile Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe
1670 1675 1680

Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser
1685 1690 1695

Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg
1700 1705 1710

Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu
1715 1720 1725

Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala
1730 1735 1740

Gly Ser Pro Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu
1745 1750 1755

Glu Asp Leu Ser Pro Ala Thr Gly His His His His His His
1760 1765 1770

<210> 988
<211> 5319
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 988
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acggggacgg gaccgggggc gtcaccagga acgtcctcca ccggctcgcc gggtgcatca 180

cccggaacgt catcgaccgg atcgccaggg agctcgacgc catcaggcgc aacaggatca 240

cctggctcaa gccctagcgc gtcaaccggc acgggtccgg gtgcctcccc tggcacgtcc 300

agcaccggat cacccggatc gagcccatcc gcctcaaccg gaaccggacc cggtacacca 360

gggtcgggaa cagcctcctc gtcaccaggc tcctcaaccc cctcgggagc cacgggttcg 420

cccggttcgt caacgccttc cggagcaact ggtagccccg gagcatcgcc aggaacttcg 480

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ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

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accttgcaca aatttatact acttttgct gtatttgatg aagggaaaag ttggcactca 1080

gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

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tacacagatg aaacctttaa gactcgtgaa gctattcagc atgaatcagg aatcttggga 1800

cctttacttt atggggaagt tggagacaca ctgttgatta tatttaagaa tcaagcaagc 1860

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ctgtcacctg caaccggtca tcatcaccat caccattga 5319

<210> 989
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 989

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Thr Ser Glu
20 25 30

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
35 40 45

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
50 55 60

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
65 70 75 80

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
85 90 95

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
100 105 110

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
115 120 125

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
130 135 140

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu

145 150 155 160

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ala Ser Ser
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
260 265 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
275 280 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
290 295 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305 310 315 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
325 330 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
340 345 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
355 360 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
370 375 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385 390 395 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
405 410 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg

420 425 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
435 440 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
450 455 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465 470 475 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
485 490 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
500 505 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
515 520 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
530 535 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545 550 555 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
565 570 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
580 585 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
595 600 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
610 615 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
645 650 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys

690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
835 840 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
885 890 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
900 905 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
915 920 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
930 935 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945 950 955 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn

965 970 975

Arg Ala Gln Ser Gly Gly Ala Pro Thr Ser Glu Ser Ala Thr Pro Glu
980 985 990

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
995 1000 1005

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
1010 1015 1020

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
1025 1030 1035

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
1040 1045 1050

Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
1055 1060 1065

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala
1070 1075 1080

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro
1085 1090 1095

Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
1100 1105 1110

Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
1115 1120 1125

Ala Ser Ser Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu
1130 1135 1140

Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
1145 1150 1155

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1160 1165 1170

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1175 1180 1185

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1190 1195 1200

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1205 1210 1215

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys

```
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
```

1475 1480 1485

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1490 1495 1500

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1505 1510 1515

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1520 1525 1530

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1535 1540 1545

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1550 1555 1560

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1565 1570 1575

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1580 1585 1590

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1595 1600 1605

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1610 1615 1620

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1625 1630 1635

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro

1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 990
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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gggagcgagc cagccacatc tgggtcggaa acgccaggca caagtgagtc tgcaactccc 180

gagtccggac ctggctccga gcctgccact agcggctccg agactccggg aacttccgag 240

agcgctacac cagaaagcgg acccggaacc agtaccgaac ctagcgaggg ctctgctccg 300

ggcagcccag ccggctctcc tacatccacg gaggagggca cttccgaatc cgccaccccg 360

gagtcagggc caggatctga acccgctacc tcaggcagtg agacgccagg aacgagcgag 420

tccgctacac cggagagtgg gccagggagc cctgctggat ctcctacgtc cactgaggaa 480

gggtcaccag cgggctcgcc caccagcact gaagaaggtg cctcgagcga cgcaagattt 540

cctcctagag tgccaaaatc ttttccattc aacacctcag tcgtgtacaa aaagactctg 600

tttgtagaat tcacggatca ccttttcaac atcgctaagc caaggccacc ctggatgggt 660

ctgctaggtc ctaccatcca ggctgaggtt tatgatacag tggtcattac acttaagaac 720

atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag 780

ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct 840

ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

ggcctcattg gagccctact agtatgtaga gaagggagtc tggccaagga aaagacacag 1020

accttgcaca aatttatact acttttgct gtatttgatg aagggaaaag ttggcactca 1080

gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg 1200

aaatcagtct attggcatgt gattggaatg ggcaccactc ctgaagtgca ctcaatattc 1260

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ataactttcc ttactgctca aacactcttg atggaccttg gacagtttct actgttttgt 1380

catatctctt cccaccaaca tgatggcatg gaagcttatg tcaaagtaga cagctgtcca 1440

gaggaacccc aactacgaat gaaaaataat gaagaagcgg aagactatga tgatgatctt 1500

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tatttgaaca atggccctca gcggattggt aggaagtaca aaaaagtccg atttatggca 1740

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cctttacttt atggggaagt tggagacaca ctgttgatta tatttaagaa tcaagcaagc 1860

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cgccatcaac gggaaataac tcgtactact cttcagtcag atcaagagga aatcgattat 2760

gatgatacca tatcagttga aatgaagaag gaagattttg acatttatga tgaggatgaa 2820

aatcagagcc cccgcagctt tcaaaagaaa acacgacact attttattgc tgcagtggag 2880

aggctctggg attatgggat gagtagctcc ccacatgttc taagaaacag ggctcagagt 2940

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acaagcggaa gcgaaactcc ggggacctca gagtctgcca ctcccgaatc ggggccaggc 3060

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gcaaccggtc atcatcacca tcaccattga 5310

<210> 991
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 991

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Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Gly Ala Pro Gly Ser Ser Pro Ser
        35                  40                  45

Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
    50                  55                  60

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65                  70                  75                  80

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro
                85                  90                  95

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
            100                 105                 110

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
        115                 120                 125

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser
    130                 135                 140

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145                 150                 155                 160

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
                165                 170                 175

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255
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Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525
```

```
Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800
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Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Gly Ala Pro Thr Ser Glu Ser Ala Thr Pro Glu
            980                 985                 990

Ser Gly Pro Gly Ser Glu Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly
        995                 1000                1005

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro
    1010                1015                1020

Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr
    1025                1030                1035

Pro Glu  Ser Gly Pro Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    1040                1045                1050

Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    1055                1060                1065
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Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala
    1070                 1075                 1080

Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro
    1085                 1090                 1095

Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1100                 1105                 1110

Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    1115                 1120                 1125

Ala Ser  Ser Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                 1135                 1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                 1150                 1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                 1165                 1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                 1180                 1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                 1195                 1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                 1210                 1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                 1225                 1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                 1240                 1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                 1255                 1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                 1270                 1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                 1285                 1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                 1300                 1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                 1315                 1320
```

```
Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575
```

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1580 1585 1590

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1595 1600 1605

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1610 1615 1620

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1625 1630 1635

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 992
<211> 5310
<212> DNA
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<223> Description of Artificial Sequence: Synthetic polynucleotide

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<210> 993
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 993

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Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
        35                  40                  45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
    50                  55                  60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65                  70                  75                  80
```

```
Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
                85                  90                  95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
            100                 105                 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
        115                 120                 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
    130                 135                 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145                 150                 155                 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350
```

```
Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620
```

```
Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895
```

```
Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Gly Ala Pro Thr Ser Glu Ser Ala Thr Pro Glu
            980                 985                 990

Ser Gly Pro Gly Ser Glu Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly
        995                 1000                1005

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro
    1010                1015                1020

Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr
    1025                1030                1035

Pro Glu  Ser Gly Pro Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    1040                1045                1050

Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    1055                1060                1065

Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala
    1070                1075                1080

Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro
    1085                1090                1095

Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1100                1105                1110

Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    1115                1120                1125

Ala Ser  Ser Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155
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```
Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
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Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410
```

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Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665
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Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
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Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                 1690                 1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                 1705                 1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                 1720                 1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1730                 1735                 1740

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
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Ser Pro  Ala Thr Gly His His  His His His His
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<211> 5310
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<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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<211> 1769
<212> PRT
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Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
35 40 45

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
50 55 60

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
65 70 75 80

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
85 90 95

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
100 105 110

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
115 120 125

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
130 135 140

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
145 150 155 160

Ser Thr Glu Glu Gly Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
165 170 175

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Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445
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Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720
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Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Gly Ala Pro Thr Ser Glu Ser Ala Thr Pro Glu
            980                 985                 990
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Ser Gly Pro Gly Ser Glu Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly
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Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro
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Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr
    1025                1030                1035


Pro Glu  Ser Gly Pro Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    1040                1045                1050


Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    1055                1060                1065


Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala
    1070                1075                1080


Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro
    1085                1090                1095


Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1100                1105                1110


Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    1115                1120                1125


Ala Ser  Ser Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140


Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155


Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170


Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185


Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200


Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215


Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230


Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245
```

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Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
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Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
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Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
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Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
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Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500
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Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1730                1735                1740

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
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aattatcgct tccatgcaat caatggctac ataatggata cactacctgg cttagtaatg 4020

gctcaggatc aaaggattcg atggtatctg ctcagcatgg gcagcaatga aaacatccat 4080

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<210> 997
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 997
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Arg Val Pro Lys Ser Phe Pro Phe Gly Ala Pro Thr Ser Glu Ser Ala
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Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu
    50                  55                  60

Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
65                  70                  75                  80

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala
                85                  90                  95

Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
            100                 105                 110

Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr
        115                 120                 125

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr
    130                 135                 140

Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
145                 150                 155                 160

Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser
                165                 170                 175

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ala Ser Ser Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
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275 280 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
290 295 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305 310 315 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
325 330 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
340 345 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
355 360 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
370 375 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385 390 395 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
405 410 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
420 425 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
435 440 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
450 455 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465 470 475 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
485 490 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
500 505 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
515 520 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
530 535 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln

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545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
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820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
835 840 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
885 890 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
900 905 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
915 920 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
930 935 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945 950 955 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
965 970 975

Arg Ala Gln Ser Gly Gly Ala Pro Thr Ser Glu Ser Ala Thr Pro Glu
980 985 990

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
995 1000 1005

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
1010 1015 1020

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
1025 1030 1035

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
1040 1045 1050

Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
1055 1060 1065

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala
1070 1075 1080

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro

1085 1090 1095

Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
1100 1105 1110

Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
1115 1120 1125

Ala Ser Ser Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu
1130 1135 1140

Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
1145 1150 1155

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1160 1165 1170

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1175 1180 1185

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1190 1195 1200

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1205 1210 1215

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1220 1225 1230

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1235 1240 1245

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1250 1255 1260

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1265 1270 1275

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1280 1285 1290

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1295 1300 1305

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1310 1315 1320

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1325 1330 1335

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met

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Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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1595 1600 1605

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
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Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1625 1630 1635

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
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Ser Pro Ala Thr Gly His His His His His His
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<211> 5310
<212> DNA
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<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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tctgggtcgg aaacgccagg cacaagtgag tctgcaactc ccgagtccgg acctggctcc 240

gagcctgcca ctagcggctc cgagactccg ggaacttccg agagcgctac accagaaagc 300

ggacccggaa ccagtaccga acctagcgag ggctctgctc cgggcagccc agccggctct 360

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<210> 999
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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20 25 30

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
35 40 45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
50 55 60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65 70 75 80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
85 90 95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
100 105 110

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Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
        115                 120                 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
    130                 135                 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145                 150                 155                 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380
```

```
Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655
```

```
Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925
```

```
Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Gly Ala Pro Thr Ser Glu Ser Ala Thr Pro Glu
            980                 985                 990

Ser Gly Pro Gly Ser Glu Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly
        995                 1000                1005

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro
    1010                1015                1020

Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr
    1025                1030                1035

Pro Glu  Ser Gly Pro Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    1040                1045                1050

Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    1055                1060                1065

Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala
    1070                1075                1080

Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro
    1085                1090                1095

Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1100                1105                1110

Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    1115                1120                1125

Ala Ser  Ser Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185
```

```
Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440
```

```
Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695
```

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

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<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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acggggacgg gaccgggggc gtcaccagga acgtcctcca ccggctcgcc gggtgcatca 180

cccggaacgt catcgaccgg atcgccaggg agctcgacgc catcaggcgc aacaggatca 240

cctggctcaa gccctagcgc gtcaaccggc acgggtccgg gtgcctcccc tggcacgtcc 300

agcaccggat cacccggatc gagcccatcc gcctcaaccg gaaccggacc cggtacacca 360

gggtcgggaa cagcctcctc gtcaccaggc tcctcaaccc cctcgggagc cacgggttcg 420

cccggttcgt caacgccttc cggagcaact ggtagccccg gagcatcgcc aggaacttcg 480

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ctgctaggtc ctaccatcca ggctgaggtt tatgatacag tggtcattac acttaagaac 720

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ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

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agaccatata acatctaccc tcacggaatc actgatgtcc gtcctttgta ttcaaggaga 1920

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gatgatacca tatcagttga aatgaagaag gaagattttg acatttatga tgaggatgaa 2820

aatcagagcc cccgcagctt tcaaaagaaa acacgacact attttattgc tgcagtggag 2880

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<210> 1001
<211> 1772
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1001

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

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His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
                405                 410                 415

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser
            420                 425                 430

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly
        435                 440                 445

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
    450                 455                 460

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala
465                 470                 475                 480
```

```
Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly
                485                 490                 495

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
            500                 505                 510

Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala
        515                 520                 525

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
    530                 535                 540

Ser Ser Thr Gly Ser Pro Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750
```

```
Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
            980                 985                 990

Ser Pro Gly Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
        995                 1000                1005

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1010                1015                1020
```

```
Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1025                 1030                 1035

Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1040                 1045                 1050

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1055                 1060                 1065

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1070                 1075                 1080

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1085                 1090                 1095

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1100                 1105                 1110

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
    1115                 1120                 1125

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1130                 1135                 1140

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1145                 1150                 1155

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1160                 1165                 1170

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1175                 1180                 1185

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1190                 1195                 1200

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1205                 1210                 1215

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1220                 1225                 1230

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1235                 1240                 1245

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1250                 1255                 1260

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1265                 1270                 1275
```

```
Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1280                1285                1290

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1295                1300                1305

Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1310                1315                1320

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1325                1330                1335

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1340                1345                1350

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1355                1360                1365

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1370                1375                1380

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1385                1390                1395

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1400                1405                1410

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1415                1420                1425

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1430                1435                1440

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1445                1450                1455

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1460                1465                1470

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1475                1480                1485

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1490                1495                1500

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1505                1510                1515

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1520                1525                1530
```

```
Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1535                1540                1545

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1550                1555                1560

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1565                1570                1575

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1580                1585                1590

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1595                1600                1605

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1610                1615                1620

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1625                1630                1635

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1640                1645                1650

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1655                1660                1665

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1670                1675                1680

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1685                1690                1695

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    1700                1705                1710

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    1715                1720                1725

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    1730                1735                1740

Gly Ser  Pro Gly Ala Glu Thr  Ala Glu Gln Lys Leu  Ile Ser Glu
    1745                1750                1755

Glu Asp  Leu Ser Pro Ala Thr  Gly His His His His  His His
    1760                1765                1770
```

<210> 1002
<211> 5319
<212> DNA

<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 1002
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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

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tatttgaaca atggccctca gcggattggt aggaagtaca aaaaagtccg atttatggca 1740

tacacagatg aaacctttaa gactcgtgaa gctattcagc atgaatcagg aatcttggga 1800

cctttacttt atggggaagt tggagacaca ctgttgatta tatttaagaa tcaagcaagc 1860

agaccatata acatctaccc tcacggaatc actgatgtcc gtcctttgta ttcaaggaga 1920

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tataaatgga cagtgactgt agaagatggg ccaactaaat cagatcctcg gtgcctgacc 2040

cgctattact ctagtttcgt taatatggag agagatctag cttcaggact cattggccct 2100

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aggctctggg attatgggat gagtagctcc ccacatgttc taagaaacag ggctcagggc 2940

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cctgtggtga actctctaga cccaccgtta ctgactcgct accttcgaat tcacccccag 5160

agttgggtgc accagattgc cctgaggatg gaggttctgg gctgcgaggc acaggacctc 5220

tacggcgccg gatcacctgg ggccgaaacg gccgaacaaa aactcatctc agaagaggat 5280

ctgtcacctg caaccggtca tcatcaccat caccattga 5319

<210> 1003
<211> 1772
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1003
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

```
Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300
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Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Gly
385                 390                 395                 400

Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro
                405                 410                 415

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
            420                 425                 430

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
        435                 440                 445

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
    450                 455                 460

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
465                 470                 475                 480

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
                485                 490                 495

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
            500                 505                 510

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
        515                 520                 525

Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    530                 535                 540

Pro Gly Ala Ser Ser Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575
```

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Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845
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Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
            980                 985                 990

Ser Pro Gly Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
        995                 1000                1005

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1010                1015                1020

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1025                1030                1035

Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1040                1045                1050

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1055                1060                1065

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1070                1075                1080

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1085                1090                1095

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1100                1105                1110
```

```
Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
    1115                 1120                 1125

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1130                 1135                 1140

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1145                 1150                 1155

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1160                 1165                 1170

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1175                 1180                 1185

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1190                 1195                 1200

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1205                 1210                 1215

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1220                 1225                 1230

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1235                 1240                 1245

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1250                 1255                 1260

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1265                 1270                 1275

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1280                 1285                 1290

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1295                 1300                 1305

Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1310                 1315                 1320

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1325                 1330                 1335

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1340                 1345                 1350

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1355                 1360                 1365
```

```
Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1370                1375                1380

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1385                1390                1395

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1400                1405                1410

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1415                1420                1425

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1430                1435                1440

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1445                1450                1455

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1460                1465                1470

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1475                1480                1485

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1490                1495                1500

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1505                1510                1515

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1520                1525                1530

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1535                1540                1545

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1550                1555                1560

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1565                1570                1575

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1580                1585                1590

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1595                1600                1605

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1610                1615                1620
```

Arg Ser Asn Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp
1625 1630 1635

Leu Gln Val Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr
1640 1645 1650

Thr Gln Gly Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu
1655 1660 1665

Phe Leu Ile Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe
1670 1675 1680

Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser
1685 1690 1695

Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg
1700 1705 1710

Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu
1715 1720 1725

Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala
1730 1735 1740

Gly Ser Pro Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu
1745 1750 1755

Glu Asp Leu Ser Pro Ala Thr Gly His His His His His His
1760 1765 1770

<210> 1004
<211> 5319
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcggc 1200

gcgccaacca gtacggagcc gtccgagggg agcgcaccag gaagcccggc tgggagcccg 1260

acttctaccg aagagggtac atctaccgaa ccaagtgaag gttcagcacc aggcacctca 1320

acagaaccct ctgagggctc ggcgcctggt acaagtgagt ccgccacccc agaatccggg 1380

cctgggacaa gcacagaacc ttcggaaggg agtgcccctg gaacatccga atcggcaacc 1440

ccagaatcag ggccaggatc tgagcccgcg acttcgggct ccgagacgcc tgggacatcc 1500

accgagccct ccgaaggatc agccccaggc accagcacgg agccctctga gggaagcgca 1560

cctggtacca gcgaaagcgc aactcccgaa tcaggtcccg gtacgagcga gtcggcgacc 1620

ccggagagcg ggccaggtgc ctcgagcctc gcccccgatg acagaagtta taaaagtcaa 1680

tatttgaaca atggccctca gcggattggt aggaagtaca aaaaagtccg atttatggca 1740

tacacagatg aaacctttaa gactcgtgaa gctattcagc atgaatcagg aatcttggga 1800

cctttacttt atggggaagt tggagacaca ctgttgatta tatttaagaa tcaagcaagc 1860

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ctgagtaaaa acaatgccat tgaaccaaga agcttctctc aaaacccacc agtcttgaaa 2700

cgccatcaac gggaaataac tcgtactact cttcagtcag atcaagagga aatcgattat 2760

gatgatacca tatcagttga aatgaagaag gaagattttg acatttatga tgaggatgaa 2820

aatcagagcc cccgcagctt tcaaaagaaa acacgacact attttattgc tgcagtggag 2880

aggctctggg attatgggat gagtagctcc ccacatgttc taagaaacag ggctcagggc 2940

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tcaagcactg gctcccctgc ctcgagcagg gctcagagtg gcagtgtccc tcagttcaag 3420

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gtggacttcc agaagacaat gaaagtcaca ggagtaacta ctcagggagt aaaatctctg 4980

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agttgggtgc accagattgc cctgaggatg gaggttctgg gctgcgaggc acaggacctc 5220

tacggcgccg gatcacctgg ggccgaaacg gccgaacaaa aactcatctc agaagaggat 5280

ctgtcacctg caaccggtca tcatcaccat caccattga 5319

<210> 1005
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1005

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn

130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Gly Ala Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly

405 410 415

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser
420 425 430

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly
435 440 445

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
450 455 460

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala
465 470 475 480

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly
485 490 495

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
500 505 510

Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala
515 520 525

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
530 535 540

Ser Ser Thr Gly Ser Pro Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545 550 555 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
565 570 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
580 585 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
595 600 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
610 615 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
645 650 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn

675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
835 840 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
885 890 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
900 905 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
915 920 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
930 935 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu

945 950 955 960

Arg Leu Trp Asp Tyr Gly Met Gly Ala Pro Thr Ser Glu Ser Ala Thr
965 970 975

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
980 985 990

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
995 1000 1005

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala
1010 1015 1020

Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
1025 1030 1035

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
1040 1045 1050

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro
1055 1060 1065

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser
1070 1075 1080

Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
1085 1090 1095

Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
1100 1105 1110

Gly Ala Ser Ser Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala
1115 1120 1125

Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu
1130 1135 1140

Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
1145 1150 1155

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1160 1165 1170

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1175 1180 1185

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1190 1195 1200

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr

1205 1210 1215

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1220 1225 1230

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1235 1240 1245

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1250 1255 1260

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1265 1270 1275

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1280 1285 1290

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1295 1300 1305

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1310 1315 1320

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1325 1330 1335

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1340 1345 1350

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1355 1360 1365

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1370 1375 1380

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1385 1390 1395

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1400 1405 1410

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1415 1420 1425

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1430 1435 1440

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1445 1450 1455

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro

1460 1465 1470

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1475 1480 1485

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1490 1495 1500

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1505 1510 1515

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1520 1525 1530

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1535 1540 1545

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1550 1555 1560

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1565 1570 1575

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1580 1585 1590

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1595 1600 1605

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1610 1615 1620

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1625 1630 1635

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu

1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 1006
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 1006
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ctcggtgagc tgcctgtgga cgcaagattt cctcctagag tgccaaaatc ttttccattc 120

aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

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gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

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tccccctcgg cctcaaccgg gacgggtccc ggctcatccc ccagcgccag cactggaaca 1320

ggtcctggca ctcctggttc cggtacggca tcgtcatccc cgggaagctc aacaccgtcc 1380

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<212> PRT
<213> Artificial Sequence

<220>
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<400> 1007

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240
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Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
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Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
                405                 410                 415

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            420                 425                 430

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
        435                 440                 445

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    450                 455                 460

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
465                 470                 475                 480

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
                485                 490                 495

Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            500                 505                 510
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Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
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Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
    530                 535                 540

Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780
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```
Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
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Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
            980                 985                 990

Ser Pro Gly Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
        995                 1000                1005

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1010                1015                1020

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
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Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
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Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
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Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
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Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
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Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1100                 1105                 1110

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
    1115                 1120                 1125

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
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Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
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Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
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Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
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Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
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Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
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Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1220                 1225                 1230

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1235                 1240                 1245

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1250                 1255                 1260

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
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Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
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Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
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Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
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Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
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Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
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Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
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Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
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Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1385                 1390                 1395

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1400                 1405                 1410

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1415                 1420                 1425

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
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Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
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Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1460                 1465                 1470

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
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Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
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Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
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Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1520                 1525                 1530

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
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Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
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His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
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Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
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Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
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Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
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Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
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Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
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Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
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Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
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Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
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Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
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Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
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Gly Ser  Pro Gly Ala Glu Thr  Ala Glu Gln Lys Leu  Ile Ser Glu
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Glu Asp  Leu Ser Pro Ala Thr  Gly His His His His  His His
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

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Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
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Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
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Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
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Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
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Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser
            420                 425                 430

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly
        435                 440                 445

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
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Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala
465                 470                 475                 480

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly
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Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
            500                 505                 510

Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala
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Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
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Ser Ser Thr Gly Ser Pro Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605
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Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
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Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
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Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
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Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
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Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
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Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Gly Ala Pro Thr Ser Glu Ser Ala Thr Pro Glu
            980                 985                 990

Ser Gly Pro Gly Ser Glu Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly
        995                 1000                1005

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro
    1010                1015                1020

Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr
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Pro Glu  Ser Gly Pro Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    1040                1045                1050

Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    1055                1060                1065

Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala
    1070                1075                1080

Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro
    1085                1090                1095

Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1100                1105                1110

Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    1115                1120                1125

Ala Ser  Ser Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
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Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
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Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
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Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
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Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
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Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
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Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
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Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
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Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
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Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
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Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
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Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
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Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
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Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
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Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
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Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
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Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
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Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
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Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
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Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
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Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
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Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
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Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
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<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1011
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

```
His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
                405                 410                 415

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            420                 425                 430
```

```
Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
        435                 440                 445

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    450                 455                 460

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
465                 470                 475                 480

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
                485                 490                 495

Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            500                 505                 510

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
        515                 520                 525

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
    530                 535                 540

Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700
```

```
Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975
```

```
Arg Ala Gln Ser Gly Gly Ala Pro Thr Ser Glu Ser Ala Thr Pro Glu
            980                 985                 990

Ser Gly Pro Gly Ser Glu Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly
        995                 1000                1005

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro
    1010                1015                1020

Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr
    1025                1030                1035

Pro Glu  Ser Gly Pro Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    1040                1045                1050

Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    1055                1060                1065

Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala
    1070                1075                1080

Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro
    1085                1090                1095

Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1100                1105                1110

Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    1115                1120                1125

Ala Ser  Ser Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230
```

```
Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485
```

```
Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1730                1735                1740
```

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 1012
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg gcgcgccaac cagtacggag ccgtccgagg ggagcgcacc aggaagcccg 1260

gctgggagcc cgacttctac cgaagagggt acatctaccg aaccaagtga aggttcagca 1320

ccaggcacct caacagaacc ctctgagggc tcggcgcctg gtacaagtga gtccgccacc 1380

ccagaatccg ggcctgggac aagcacagaa ccttcggaag ggagtgcccc tggaacatcc 1440

gaatcggcaa ccccagaatc agggccagga tctgagcccg cgacttcggg ctccgagacg 1500

cctgggacat ccaccgagcc ctccgaagga tcagccccag gcaccagcac ggagccctct 1560

gagggaagcg cacctggtac cagcgaaagc gcaactcccg aatcaggtcc cggtacgagc 1620

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tacacagatg aaacctttaa gactcgtgaa gctattcagc atgaatcagg aatcttggga 1800

cctttacttt atggggaagt tggagacaca ctgttgatta tatttaagaa tcaagcaagc 1860

agaccatata acatctaccc tcacggaatc actgatgtcc gtcctttgta ttcaaggaga 1920

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tataaatgga cagtgactgt agaagatggg ccaactaaat cagatcctcg gtgcctgacc 2040

cgctattact ctagtttcgt taatatggag agagatctag cttcaggact cattggccct 2100

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tccaacatca tgcacagcat caatggctat gtttttgata gtttgcagtt gtcagtttgt 2340

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gtcttcttct ctggatatac cttcaaacac aaaatggtct atgaagacac actcacccta 2460

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cgccatcaac gggaaataac tcgtactact cttcagtcag atcaagagga aatcgattat 2760

gatgatacca tatcagttga aatgaagaag gaagattttg acatttatga tgaggatgaa 2820

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aggctctggg attatgggat gagtagctcc ccacatgttc taagaaacag ggctcagagt 2940

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acaagcggaa gcgaaactcc ggggacctca gagtctgcca ctcccgaatc ggggccaggc 3060

tctgaaccgg ccacttcagg gagcgaaaca ccaggaacat cggagagcgc taccccggag 3120

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tgggcttatt tctctgatgt tgacctggaa aaagatgtgc actcaggcct gattggaccc 3780

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atggaaagaa actgcagggc tccctgcaat atccagatgg aagatcccac ttttaaagag 3960

aattatcgct tccatgcaat caatggctac ataatggata cactacctgg cttagtaatg 4020

gctcaggatc aaaggattcg atggtatctg ctcagcatgg gcagcaatga aaacatccat 4080

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atttggcggg tggaatgcct tattggcgag catctacatg ctgggatgag cacacttttt 4260

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ataaaacaca atatttttaa ccctccaatt attgctcgat acatccgttt gcacccaact 4680

cattatagca ttcgcagcac tcttcgcatg gagttgatgg gctgtgattt aaatagttgc 4740

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gcaaccggtc atcatcacca tcaccattga 5310

<210> 1013
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic

polypeptide

<400> 1013

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Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1               5                  10                  15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
```

260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Gly
385 390 395 400

Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro
405 410 415

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
420 425 430

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
435 440 445

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
450 455 460

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
465 470 475 480

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
485 490 495

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
500 505 510

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
515 520 525

Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly

530 535 540

Pro Gly Ala Ser Ser Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545 550 555 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
565 570 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
580 585 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
595 600 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
610 615 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
645 650 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp

805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
835 840 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
885 890 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
900 905 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
915 920 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
930 935 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945 950 955 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
965 970 975

Arg Ala Gln Ser Gly Gly Ala Pro Thr Ser Glu Ser Ala Thr Pro Glu
980 985 990

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
995 1000 1005

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
1010 1015 1020

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
1025 1030 1035

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
1040 1045 1050

Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
1055 1060 1065

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala

1070 1075 1080

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro
1085 1090 1095

Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
1100 1105 1110

Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
1115 1120 1125

Ala Ser Ser Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu
1130 1135 1140

Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
1145 1150 1155

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1160 1165 1170

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1175 1180 1185

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1190 1195 1200

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1205 1210 1215

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1220 1225 1230

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1235 1240 1245

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1250 1255 1260

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1265 1270 1275

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1280 1285 1290

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1295 1300 1305

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1310 1315 1320

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu

1325 1330 1335

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1340 1345 1350

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1355 1360 1365

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1370 1375 1380

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1385 1390 1395

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1400 1405 1410

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1415 1420 1425

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1430 1435 1440

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1445 1450 1455

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1460 1465 1470

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1475 1480 1485

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1490 1495 1500

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1505 1510 1515

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1520 1525 1530

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1535 1540 1545

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1550 1555 1560

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1565 1570 1575

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp

1580 1585 1590

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1595 1600 1605

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1610 1615 1620

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1625 1630 1635

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 1014
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

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agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420
ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480
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gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600
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tactttacca atatgtttgc cacctggtct ccttcaaaag ctcgacttca cctccaaggg 4860

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cagaagacaa tgaaagtcac aggagtaact actcagggag taaaatctct gcttaccagc 4980

atgtatgtga aggagttcct catctccagc agtcaagatg gccatcagtg gactctcttt 5040

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aactctctag acccaccgtt actgactcgc taccttcgaa ttcaccccca gagttgggtg 5160

caccagattg ccctgaggat ggaggttctg ggctgcgagg cacaggacct ctacggcgcc 5220

ggatcacctg gggccgaaac ggccgaacaa aaactcatct cagaagagga tctgtcacct 5280

gcaaccggtc atcatcacca tcaccattga 5310

<210> 1015
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1015
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

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Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365
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Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
                405                 410                 415

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            420                 425                 430

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
        435                 440                 445

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    450                 455                 460

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
465                 470                 475                 480

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
                485                 490                 495

Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            500                 505                 510

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
        515                 520                 525

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
    530                 535                 540

Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640
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Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910
```

```
Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Gly Ala Pro Thr Ser Glu Ser Ala Thr
                965                 970                 975

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
            980                 985                 990

Pro Gly Thr Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
        995                 1000                1005

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1010                1015                1020

Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1040                1045                1050

Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr Glu Glu  Gly Ser Pro
    1055                1060                1065

Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Ser Pro  Ala Gly Ser
    1070                1075                1080

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1085                1090                1095

Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1100                1105                1110

Gly Ala  Ser Ser Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170
```

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Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425
```

```
Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680
```

```
Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                 1690                 1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                 1705                 1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                 1720                 1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1730                 1735                 1740

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
    1745                 1750                 1755

Ser Pro  Ala Thr Gly His His  His His His His
    1760                 1765
```

<210> 1016
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg gcgcgccaac cagtacggag ccgtccgagg ggagcgcacc aggaagcccg 1260

gctgggagcc cgacttctac cgaagagggt acatctaccg aaccaagtga aggttcagca 1320

ccaggcacct caacagaacc ctctgagggc tcggcgcctg gtacaagtga gtccgccacc 1380

ccagaatccg ggcctgggac aagcacagaa ccttcggaag ggagtgcccc tggaacatcc 1440

gaatcggcaa ccccagaatc agggccagga tctgagcccg cgacttcggg ctccgagacg 1500

cctgggacat ccaccgagcc ctccgaagga tcagccccag gcaccagcac ggagccctct 1560

gagggaagcg cacctggtac cagcgaaagc gcaactcccg aatcaggtcc cggtacgagc 1620

gagtcggcga ccccggagag cgggccaggt gcctcgagcg acagaagtta taaaagtcaa 1680

tatttgaaca atggccctca gcggattggt aggaagtaca aaaaagtccg atttatggca 1740

tacacagatg aaacctttaa gactcgtgaa gctattcagc atgaatcagg aatcttggga 1800

cctttacttt atggggaagt tggagacaca ctgttgatta tatttaagaa tcaagcaagc 1860

agaccatata acatctaccc tcacggaatc actgatgtcc gtcctttgta ttcaaggaga 1920

ttaccaaaag gtgtaaaaca tttgaaggat tttccaattc tgccaggaga aatattcaaa 1980

tataaatgga cagtgactgt agaagatggg ccaactaaat cagatcctcg gtgcctgacc 2040

cgctattact ctagtttcgt taatatggag agagatctag cttcaggact cattggccct 2100

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gggtgccaca actcagactt tcggaacaga ggcatgaccg ccttactgaa ggtttctagt 2580

tgtgacaaga acactggtga ttattacgag gacagttatg aagatatttc agcatacttg 2640

ctgagtaaaa acaatgccat tgaaccaaga agcttctctc aaaacccacc agtcttgaaa 2700

cgccatcaac gggaaataac tcgtactact cttcagtcag atcaagagga aatcgattat 2760

gatgatacca tatcagttga aatgaagaag gaagattttg acatttatga tgaggatgaa 2820

aatcagagcc cccgcagctt tcaaaagaaa acacgacact attttattgc tgcagtggag 2880

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cctggtagtg agcctgccac aagcggaagc gaaactccgg ggacctcaga gtctgccact 3000

cccgaatcgg ggccaggctc tgaaccggcc acttcaggga gcgaaacacc aggaacatcg 3060

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cccggtacct ccactgagcc cagcgagggg agtgcaccag gtgcctcgag cagtagctcc 3360

ccacatgttc taagaaacag ggctcagagt ggcagtgtcc ctcagttcaa gaaagttgtt 3420

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catttgggac tcctggggcc atatataaga gcagaagttg aagataatat catggtaact 3540

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gatcagaggc aaggagcaga acctagaaaa aactttgtca agcctaatga aaccaaaact 3660

tacttttgga aagtgcaaca tcatatggca cccactaaag atgagtttga ctgcaaagcc 3720

tgggcttatt tctctgatgt tgacctggaa aaagatgtgc actcaggcct gattggaccc 3780

cttctggtct gccacactaa cacactgaac cctgctcatg ggagacaagt gacagtacag 3840

gaatttgctc tgtttttcac catctttgat gagaccaaaa gctggtactt cactgaaaat 3900

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atttggcggg tggaatgcct tattggcgag catctacatg ctgggatgag cacacttttt 4260

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ataaaacaca atatttttaa ccctccaatt attgctcgat acatccgttt gcacccaact 4680

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tactttacca atatgtttgc cacctggtct ccttcaaaag ctcgacttca cctccaaggg 4860

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gcaaccggtc atcatcacca tcaccattga 5310

<210> 1017
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1017

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

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His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Gly
385                 390                 395                 400

Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro
                405                 410                 415

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
            420                 425                 430

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
        435                 440                 445

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
    450                 455                 460
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Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
465                 470                 475                 480

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
                485                 490                 495

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
            500                 505                 510

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
        515                 520                 525

Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    530                 535                 540

Pro Gly Ala Ser Ser Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735
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Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005
```

```
Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1145                1150                1155

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala
    1160                1165                1170

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1175                1180                1185

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1190                1195                1200

Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
    1205                1210                1215

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala
    1220                1225                1230

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1235                1240                1245

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1250                1255                1260
```

```
Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1265                 1270                 1275

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ala
    1280                 1285                 1290

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Ala Ser  Ser Ala Pro
    1295                 1300                 1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                 1315                 1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                 1330                 1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                 1345                 1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                 1360                 1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                 1375                 1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                 1390                 1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                 1405                 1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                 1420                 1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                 1435                 1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                 1450                 1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                 1465                 1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                 1480                 1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                 1495                 1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                 1510                 1515
```

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Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1730                1735                1740

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
    1745                1750                1755

Ser Pro  Ala Thr Gly His His  His His His His His
    1760                1765
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<210> 1018
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

<400> 1018
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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285
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Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
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Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
                405                 410                 415

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser
            420                 425                 430

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly
        435                 440                 445

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
    450                 455                 460

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala
465                 470                 475                 480

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly
                485                 490                 495

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
            500                 505                 510

Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala
        515                 520                 525

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
    530                 535                 540

Ser Ser Thr Gly Ser Pro Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560
```

```
Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830
```

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Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
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Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
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Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095
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Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
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Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
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Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
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Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1145                1150                1155

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala
    1160                1165                1170

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1175                1180                1185

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1190                1195                1200

Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
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Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala
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Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1235                1240                1245

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
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Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
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Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ala
    1280                1285                1290

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Ala Ser  Ser Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350
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Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
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Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
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Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
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Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
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Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
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Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
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Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val

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Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
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Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
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Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser
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Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly
435 440 445

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
450 455 460

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala
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Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly
485 490 495

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Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala
515 520 525

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
530 535 540

Ser Ser Thr Gly Ser Pro Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545 550 555 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
565 570 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
580 585 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
595 600 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
610 615 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
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Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr

660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
835 840 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
885 890 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
900 905 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
915 920 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro

930 935 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945 950 955 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
965 970 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
980 985 990

Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
995 1000 1005

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1010 1015 1020

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1025 1030 1035

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1040 1045 1050

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1055 1060 1065

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1070 1075 1080

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1085 1090 1095

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1100 1105 1110

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1115 1120 1125

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1130 1135 1140

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1145 1150 1155

Cys Asn Ile Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
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Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
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Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser

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Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser
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Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
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Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
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Ala Ser  Ser Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
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Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
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Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
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Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
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Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
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Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
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Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
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Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
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Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
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Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1520 1525 1530

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1535 1540 1545

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
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Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1565 1570 1575

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
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Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
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Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
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Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
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Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
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Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg

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Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
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aggagtaatg cctggagacc tcaggtgaat aatccaaaag agtggctgca agtggacttc 4920

cagaagacaa tgaaagtcac aggagtaact actcagggag taaaatctct gcttaccagc 4980

atgtatgtga aggagttcct catctccagc agtcaagatg gccatcagtg gactctcttt 5040

tttcagaatg gcaaagtaaa ggtttttcag ggaaatcaag actccttcac acctgtggtg 5100

aactctctag acccaccgtt actgactcgc taccttcgaa ttcaccccca gagttgggtg 5160

caccagattg ccctgaggat ggaggttctg ggctgcgagg cacaggacct ctacggcgcc 5220

ggatcacctg gggccgaaac ggccgaacaa aaactcatct cagaagagga tctgtcacct 5280

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<210> 1023
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1023
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

```
Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Gly
385                 390                 395                 400

Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro
                405                 410                 415

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
            420                 425                 430

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
        435                 440                 445

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
    450                 455                 460

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
465                 470                 475                 480

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
                485                 490                 495
```

```
Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
            500                 505                 510

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
        515                 520                 525

Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    530                 535                 540

Pro Gly Ala Ser Ser Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765
```

```
Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035
```

```
Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1145                1150                1155

Pro Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    1160                1165                1170

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1175                1180                1185

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1190                1195                1200

Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1205                1210                1215

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1220                1225                1230

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Pro  Ala Gly Ser
    1235                1240                1245

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1250                1255                1260

Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1265                1270                1275

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1280                1285                1290
```

```
Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ala Ser  Ser Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545
```

```
Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1730                1735                1740

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
    1745                1750                1755

Ser Pro  Ala Thr Gly His His  His His His His His
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<210> 1024
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcggc 1200

gcgccaacca gtacggagcc gtccgagggg agcgcaccag gaagcccggc tgggagcccg 1260

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cctgggacaa gcacagaacc ttcggaaggg agtgcccctg gaacatccga atcggcaacc 1440

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accgagccct ccgaaggatc agccccaggc accagcacgg agccctctga gggaagcgca 1560

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tatttgaaca atggccctca gcggattggt aggaagtaca aaaaagtccg atttatggca 1740

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cctttacttt atggggaagt tggagacaca ctgttgatta tatttaagaa tcaagcaagc 1860

agaccatata acatctaccc tcacggaatc actgatgtcc gtcctttgta ttcaaggaga 1920

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cgctattact ctagtttcgt taatatggag agagatctag cttcaggact cattggccct 2100

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cagcccttat accgtggaga actaaatgaa catttgggac tcctggggcc atatataaga 3060

gcagaagttg aagataatat catggtaact ttcagaaatc aggcctctcg tccctattcc 3120

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ggccctggct ctccagcggg ctcacccacg agcactgaag agggctctcc cgctggcagc 3780

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gcaccaggtg cctcgagcgc tccctgcaat atccagatgg aagatcccac ttttaaagag 3960

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<210> 1025
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

```
Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320
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```
Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
                405                 410                 415

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser
            420                 425                 430

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly
        435                 440                 445

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
    450                 455                 460

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala
465                 470                 475                 480

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly
                485                 490                 495

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
            500                 505                 510

Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala
        515                 520                 525

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
    530                 535                 540

Ser Ser Thr Gly Ser Pro Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590
```

```
Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860
```

```
Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125
```

```
Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1145                1150                1155

Pro Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    1160                1165                1170

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1175                1180                1185

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1190                1195                1200

Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1205                1210                1215

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1220                1225                1230

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Pro  Ala Gly Ser
    1235                1240                1245

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1250                1255                1260

Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1265                1270                1275

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1280                1285                1290

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ala Ser  Ser Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380
```

```
Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635
```

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 1026
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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ctcggtgagc tgcctgtgga cgcaagattt cctcctagag tgccaaaatc ttttccattc 120

aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg gcgcgccagg tgcctcgccg ggaacatcat caactggttc acccgggtca 1260

tccccctcgg cctcaaccgg gacgggtccc ggctcatccc ccagcgccag cactggaaca 1320

ggtcctggca ctcctggttc cggtacggca tcgtcatccc cgggaagctc aacaccgtcc 1380

ggagcgacag gatcacctgg ctcgtcacct tcggcgtcaa ctggaacggg gccaggggcc 1440

tcacccggaa cgtcctcgac tgggtcgcct ggtacgccgg gatcaggaac ggcctcatcc 1500

tcgcctgggt cctcaacgcc ctcgggtgcg actggttcgc cgggaactcc tggctcgggg 1560

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tatttgaaca atggccctca gcggattggt aggaagtaca aaaaagtccg atttatggca 1740

tacacagatg aaacctttaa gactcgtgaa gctattcagc atgaatcagg aatcttggga 1800

cctttacttt atggggaagt tggagacaca ctgttgatta tatttaagaa tcaagcaagc 1860

agaccatata acatctaccc tcacggaatc actgatgtcc gtcctttgta ttcaaggaga 1920

ttaccaaaag gtgtaaaaca tttgaaggat tttccaattc tgccaggaga aatattcaaa 1980

tataaatgga cagtgactgt agaagatggg ccaactaaat cagatcctcg gtgcctgacc 2040

cgctattact ctagtttcgt taatatggag agagatctag cttcaggact cattggccct 2100

ctcctcatct gctacaaaga atctgtagat caaagaggaa accagataat gtcagacaag 2160

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tccaacatca tgcacagcat caatggctat gtttttgata gtttgcagtt gtcagtttgt 2340

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gggtgccaca actcagactt tcggaacaga ggcatgaccg ccttactgaa ggtttctagt 2580

tgtgacaaga acactggtga ttattacgag gacagttatg aagatatttc agcatacttg 2640

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cgccatcaac gggaaataac tcgtactact cttcagtcag atcaagagga aatcgattat 2760

gatgatacca tatcagttga aatgaagaag gaagattttg acatttatga tgaggatgaa 2820

aatcagagcc cccgcagctt tcaaaagaaa acacgacact attttattgc tgcagtggag 2880

aggctctggg attatgggat gagtagctcc ccacatgttc taagaaacag ggctcagagt 2940

ggcagtgtcc ctcagttcaa gaaagttgtt ttccaggaat ttactgatgg ctcctttact 3000

cagcccttat accgtggaga actaaatgaa catttgggac tcctggggcc atatataaga 3060

gcagaagttg aagataatat catggtaact ttcagaaatc aggcctctcg tccctattcc 3120

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tcagggagcg aaacaccagg aacatcggag agcgctaccc cggagagcgg gccaggaact 3660

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ggccctggct ctccagcggg ctcacccacg agcactgaag agggctctcc cgctggcagc 3780

ccaacgtcga cagaagaagg atcaccagca ggctccccca catcaacaga ggagggtaca 3840

tcagaatctg ctactcccga gagtggaccc ggtacctcca ctgagcccag cgaggggagt 3900

gcaccaggtg cctcgagcgc tccctgcaat atccagatgg aagatcccac ttttaaagag 3960

aattatcgct tccatgcaat caatggctac ataatggata cactacctgg cttagtaatg 4020

gctcaggatc aaaggattcg atggtatctg ctcagcatgg gcagcaatga aaacatccat 4080

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ctgtacaatc tctatccagg tgtttttgag acagtggaaa tgttaccatc caaagctgga 4200

atttggcggg tggaatgcct tattggcgag catctacatg ctgggatgag cacacttttt 4260

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tactttacca atatgtttgc cacctggtct ccttcaaaag ctcgacttca cctccaaggg 4860

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cagaagacaa tgaaagtcac aggagtaact actcagggag taaaatctct gcttaccagc 4980

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<210> 1027
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1027
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

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Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
                405                 410                 415
```

```
Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            420                 425                 430

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
        435                 440                 445

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    450                 455                 460

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
465                 470                 475                 480

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
                485                 490                 495

Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            500                 505                 510

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
        515                 520                 525

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
    530                 535                 540

Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685
```

```
Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960
```

```
Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1145                1150                1155

Pro Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    1160                1165                1170

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1175                1180                1185

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1190                1195                1200

Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1205                1210                1215
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Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1220                 1225                 1230

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Pro  Ala Gly Ser
    1235                 1240                 1245

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1250                 1255                 1260

Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1265                 1270                 1275

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1280                 1285                 1290

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ala Ser  Ser Ala Pro
    1295                 1300                 1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                 1315                 1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                 1330                 1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                 1345                 1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                 1360                 1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                 1375                 1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                 1390                 1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                 1405                 1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                 1420                 1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                 1435                 1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                 1450                 1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                 1465                 1470
```

```
Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725
```

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 1028
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg gcgcgccaac cagtacggag ccgtccgagg ggagcgcacc aggaagcccg 1260

gctgggagcc cgacttctac cgaagagggt acatctaccg aaccaagtga aggttcagca 1320

ccaggcacct caacagaacc ctctgagggc tcggcgcctg gtacaagtga gtccgccacc 1380

ccagaatccg ggcctgggac aagcacagaa ccttcggaag ggagtgcccc tggaacatcc 1440

gaatcggcaa ccccagaatc agggccagga tctgagcccg cgacttcggg ctccgagacg 1500

cctgggacat ccaccgagcc ctccgaagga tcagccccag gcaccagcac ggagccctct 1560

gagggaagcg cacctggtac cagcgaaagc gcaactcccg aatcaggtcc cggtacgagc 1620

gagtcggcga ccccggagag cgggccaggt gcctcgagcg acagaagtta taaaagtcaa 1680

tatttgaaca atggccctca gcggattggt aggaagtaca aaaaagtccg atttatggca 1740

tacacagatg aaacctttaa gactcgtgaa gctattcagc atgaatcagg aatcttggga 1800

cctttacttt atggggaagt tggagacaca ctgttgatta tatttaagaa tcaagcaagc 1860

agaccatata acatctaccc tcacggaatc actgatgtcc gtcctttgta ttcaaggaga 1920

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tataaatgga cagtgactgt agaagatggg ccaactaaat cagatcctcg gtgcctgacc 2040

cgctattact ctagtttcgt taatatggag agagatctag cttcaggact cattggccct 2100

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ctgagtaaaa acaatgccat tgaaccaaga agcttctctc aaaacccacc agtcttgaaa 2700

cgccatcaac gggaaataac tcgtactact cttcagtcag atcaagagga aatcgattat 2760

gatgatacca tatcagttga aatgaagaag gaagattttg acatttatga tgaggatgaa 2820

aatcagagcc cccgcagctt tcaaaagaaa acacgacact attttattgc tgcagtggag 2880

aggctctggg attatgggat gagtagctcc ccacatgttc taagaaacag ggctcagagt 2940

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cagcccttat accgtggaga actaaatgaa catttgggac tcctggggcc atatataaga 3060

gcagaagttg aagataatat catggtaact ttcagaaatc aggcctctcg tccctattcc 3120

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actccgggga cctcagagtc tgccactccc gaatcggggc caggctctga accggccact 3600

tcagggagcg aaacaccagg aacatcggag agcgctaccc cggagagcgg gccaggaact 3660

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ataaaacaca atatttttaa ccctccaatt attgctcgat acatccgttt gcacccaact 4680

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ggatcacctg gggccgaaac ggccgaacaa aaactcatct cagaagagga tctgtcacct 5280

gcaaccggtc atcatcacca tcaccattga 5310

<210> 1029
<211> 1769
<212> PRT

<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 1029
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His

245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Gly
385 390 395 400

Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro
405 410 415

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
420 425 430

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
435 440 445

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
450 455 460

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
465 470 475 480

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
485 490 495

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
500 505 510

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr

515 520 525

Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
530 535 540

Pro Gly Ala Ser Ser Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545 550 555 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
565 570 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
580 585 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
595 600 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
610 615 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
645 650 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser

785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
835 840 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
885 890 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
900 905 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
915 920 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
930 935 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945 950 955 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
965 970 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
980 985 990

Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
995 1000 1005

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1010 1015 1020

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1025 1030 1035

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1040 1045 1050

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr

1055 1060 1065

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1070 1075 1080

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1085 1090 1095

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1100 1105 1110

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1115 1120 1125

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1130 1135 1140

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1145 1150 1155

Cys Asn Ile Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
1160 1165 1170

Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
1175 1180 1185

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
1190 1195 1200

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala
1205 1210 1215

Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
1220 1225 1230

Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
1235 1240 1245

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
1250 1255 1260

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser
1265 1270 1275

Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
1280 1285 1290

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
1295 1300 1305

Ala Ser Ser Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg

1310 1315 1320

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1325 1330 1335

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1340 1345 1350

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1355 1360 1365

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1370 1375 1380

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1385 1390 1395

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1400 1405 1410

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1415 1420 1425

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1430 1435 1440

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1445 1450 1455

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1460 1465 1470

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1475 1480 1485

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1490 1495 1500

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1505 1510 1515

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1520 1525 1530

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1535 1540 1545

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1550 1555 1560

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn

1565 1570 1575

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1580 1585 1590

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1595 1600 1605

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1610 1615 1620

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1625 1630 1635

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 1030
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

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Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350
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Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
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Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
                405                 410                 415

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            420                 425                 430

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
        435                 440                 445

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    450                 455                 460

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
465                 470                 475                 480

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
                485                 490                 495

Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            500                 505                 510

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
        515                 520                 525

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
    530                 535                 540

Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620
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Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895
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Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
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Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
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Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155
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Cys Asn  Ile Gly Ala Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
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Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
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Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1190                 1195                 1200

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala
    1205                 1210                 1215

Ser Thr  Gly Thr Gly Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    1220                 1225                 1230

Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1235                 1240                 1245

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1250                 1255                 1260

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser
    1265                 1270                 1275

Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    1280                 1285                 1290

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1295                 1300                 1305

Ala Ser  Ser Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                 1315                 1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                 1330                 1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                 1345                 1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                 1360                 1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                 1375                 1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                 1390                 1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                 1405                 1410
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Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
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Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
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Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
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Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
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1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
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Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
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<212> PRT
<213> Artificial Sequence

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

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Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
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His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
                405                 410                 415

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser
            420                 425                 430

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly
        435                 440                 445
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Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
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Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala
465                 470                 475                 480

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly
                485                 490                 495

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
            500                 505                 510

Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala
        515                 520                 525

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
    530                 535                 540

Ser Ser Thr Gly Ser Pro Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720
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Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990
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Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
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Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
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Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Gly Ala Pro Gly  Thr Pro Gly
    1160                1165                1170

Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1175                1180                1185

Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1190                1195                1200

Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly
    1205                1210                1215

Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly Pro Gly  Thr Pro Gly
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Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
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Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
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Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly
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Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr
    1280                 1285                 1290

Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1295                 1300                 1305

Ser Thr  Gly Ser Pro Ala Ser  Ser Thr Phe Lys Glu  Asn Tyr Arg
    1310                 1315                 1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                 1330                 1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                 1345                 1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                 1360                 1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                 1375                 1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                 1390                 1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
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Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
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Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
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Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                 1450                 1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                 1465                 1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                 1480                 1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                 1495                 1500
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Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
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Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
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Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
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Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1730                1735                1740

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
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<210> 1035
<211> 1919
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1035

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            20                  25                  30

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
        35                  40                  45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
    50                  55                  60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65                  70                  75                  80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
                85                  90                  95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
            100                 105                 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
        115                 120                 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
    130                 135                 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145                 150                 155                 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270
```

```
Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
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Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540
```

```
Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu
545                 550                 555                 560

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
                565                 570                 575

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            580                 585                 590

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
        595                 600                 605

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    610                 615                 620

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
625                 630                 635                 640

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
                645                 650                 655

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
            660                 665                 670

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
        675                 680                 685

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp
    690                 695                 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815
```

```
Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020

Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Gly Ala Pro  Gly Thr Pro
    1055                1060                1065

Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly Ala Ser  Pro Gly Thr
    1070                1075                1080
```

```
Ser Ser  Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
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Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1100                1105                1110

Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr Gly Pro  Gly Thr Pro
    1115                1120                1125

Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly Ala Ser  Pro Gly Thr
    1130                1135                1140

Ser Ser  Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
    1145                1150                1155

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1160                1165                1170

Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser
    1175                1180                1185

Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    1190                1195                1200

Ser Ser  Thr Gly Ser Pro Ala  Ser Ser Ser Asp Gln  Glu Glu Ile
    1205                1210                1215

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1220                1225                1230

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1235                1240                1245

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1250                1255                1260

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1265                1270                1275

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1280                1285                1290

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1295                1300                1305

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1310                1315                1320

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1325                1330                1335
```

```
Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
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Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1355                1360                1365

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1370                1375                1380

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1385                1390                1395

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1400                1405                1410

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1415                1420                1425

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1430                1435                1440

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1445                1450                1455

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                1555                1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
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```

```
Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
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Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                1615                1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845
```

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1850 1855 1860

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1865 1870 1875

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
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Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
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actcagggag taaaatctct gcttaccagc atgtatgtga aggagttcct catctccagc 5460

agtcaagatg gccatcagtg gactctcttt tttcagaatg gcaaagtaaa ggtttttcag 5520

ggaaatcaag actccttcac acctgtggtg aactctctag acccaccgtt actgactcgc 5580

taccttcgaa ttcaccccca gagttgggtg caccagattg ccctgaggat ggaggttctg 5640

ggctgcgagg cacaggacct ctacggcgcc ggatcacctg gggccgaaac ggccgaacaa 5700

aaactcatct cagaagagga tctgtcacct gcaaccggtc atcatcacca tcaccattga 5760

<210> 1037
<211> 1922
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1037

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Gly Ser Ser
20 25 30

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
35 40 45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
50 55 60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65 70 75 80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
85 90 95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
100 105 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
115 120 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
130 135 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145 150 155 160

```
Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430
```

```
Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu
545                 550                 555                 560

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
                565                 570                 575

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            580                 585                 590

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
        595                 600                 605

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    610                 615                 620

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
625                 630                 635                 640

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
                645                 650                 655

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
            660                 665                 670

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
        675                 680                 685

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp
    690                 695                 700
```

```
Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975
```

```
Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020

Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Gly  Ala Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1130                1135                1140

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1145                1150                1155

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1160                1165                1170

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1175                1180                1185

Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1190                1195                1200

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1205                1210                1215

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1220                1225                1230
```

```
Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1235                 1240                 1245

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1250                 1255                 1260

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
    1265                 1270                 1275

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1280                 1285                 1290

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1295                 1300                 1305

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1310                 1315                 1320

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1325                 1330                 1335

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1340                 1345                 1350

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1355                 1360                 1365

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1370                 1375                 1380

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1385                 1390                 1395

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1400                 1405                 1410

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1415                 1420                 1425

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1430                 1435                 1440

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1445                 1450                 1455

Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1460                 1465                 1470

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1475                 1480                 1485
```

```
Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1490                1495                1500

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1505                1510                1515

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1520                1525                1530

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1535                1540                1545

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1550                1555                1560

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1565                1570                1575

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1580                1585                1590

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1595                1600                1605

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1610                1615                1620

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1625                1630                1635

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1640                1645                1650

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1655                1660                1665

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1670                1675                1680

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1685                1690                1695

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1700                1705                1710

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1715                1720                1725

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1730                1735                1740
```

```
Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1745                1750                1755

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1760                1765                1770

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1775                1780                1785

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1790                1795                1800

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1805                1810                1815

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1820                1825                1830

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1835                1840                1845

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    1850                1855                1860

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    1865                1870                1875

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    1880                1885                1890

Gly Ser  Pro Gly Ala Glu Thr  Ala Glu Gln Lys Leu  Ile Ser Glu
    1895                1900                1905

Glu Asp  Leu Ser Pro Ala Thr  Gly His His His His  His His
    1910                1915                1920
```

<210> 1038
<211> 5769
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 1038

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ccaggttcgt caccctcagc atcaacgggg acgggaccgg gggcgtcacc aggaacgtcc      180

tccaccggct cgccgggtgc atcacccgga acgtcatcga ccggatcgcc agggagctcg      240

acgccatcag gcgcaacagg atcacctggc tcaagcccta gcgcgtcaac cggcacgggt      300
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ccgggtgcct cccctggcac gtccagcacc ggatcacccg gatcgagccc atccgcctca 360
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acccccctcgg gagccacggg ttcgcccggt tcgtcaacgc cttccggagc aactggtagc 480
cccggagcat cgccaggaac ttcgagcacg gggtcgcccg cctcgagcga cgcaagattt 540
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actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 2400

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<210> 1039
<211> 1919
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1039
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Gly Ser Ser
20 25 30

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Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
        35                  40                  45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
    50                  55                  60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65                  70                  75                  80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
                85                  90                  95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
            100                 105                 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
        115                 120                 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
    130                 135                 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145                 150                 155                 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300
```

```
Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu
545                 550                 555                 560

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
                565                 570                 575
```

```
Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
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Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
        595                 600                 605

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    610                 615                 620

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
625                 630                 635                 640

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
                645                 650                 655

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
            660                 665                 670

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
        675                 680                 685

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp
    690                 695                 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845
```

```
Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
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Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020

Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110
```

```
Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
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Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                 1135                 1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                 1150                 1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                 1165                 1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                 1180                 1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                 1195                 1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                 1210                 1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                 1225                 1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                 1240                 1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                 1255                 1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                 1270                 1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                 1285                 1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1295                 1300                 1305

Pro Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    1310                 1315                 1320

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1325                 1330                 1335

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1340                 1345                 1350

Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1355                 1360                 1365
```

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Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1370                1375                1380

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Pro  Ala Gly Ser
    1385                1390                1395

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1400                1405                1410

Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1415                1420                1425

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1430                1435                1440

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ala Ser  Ser Ala Pro
    1445                1450                1455

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                1555                1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                1615                1620
```

```
Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875
```

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1880 1885 1890

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1895 1900 1905

Ser Pro Ala Thr Gly His His His His His His
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tccaccggct cgccgggtgc atcacccgga acgtcatcga ccggatcgcc agggagctcg 240

acgccatcag gcgcaacagg atcacctggc tcaagcccta gcgcgtcaac cggcacgggt 300

ccgggtgcct cccctggcac gtccagcacc ggatcacccg gatcgagccc atccgcctca 360

accggaaccg gacccggtac accagggtcg ggaacagcct cctcgtcacc aggctcctca 420

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Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
35 40 45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
50 55 60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65 70 75 80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
85 90 95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
100 105 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
115 120 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
130 135 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145 150 155 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

```
Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460
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```
His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735
```

```
Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
        915                 920                 925

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
    930                 935                 940

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
945                 950                 955                 960

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
                965                 970                 975

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly
            980                 985                 990

Thr Ser Ser Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
        995                 1000                1005
```

```
Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1010                 1015                 1020

Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser
    1025                 1030                 1035

Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    1040                 1045                 1050

Ser Ser  Thr Gly Ser Pro Ala  Ser Ser Ser Asp Gln  Glu Glu Ile
    1055                 1060                 1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                 1075                 1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                 1090                 1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                 1105                 1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                 1120                 1125

Gln Gly  Ala Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1130                 1135                 1140

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1145                 1150                 1155

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1160                 1165                 1170

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1175                 1180                 1185

Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1190                 1195                 1200

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1205                 1210                 1215

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1220                 1225                 1230

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1235                 1240                 1245

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1250                 1255                 1260
```

```
Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
    1265                 1270                 1275

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1280                 1285                 1290

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1295                 1300                 1305

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1310                 1315                 1320

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1325                 1330                 1335

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1340                 1345                 1350

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1355                 1360                 1365

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1370                 1375                 1380

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1385                 1390                 1395

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1400                 1405                 1410

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1415                 1420                 1425

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1430                 1435                 1440

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1445                 1450                 1455

Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1460                 1465                 1470

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1475                 1480                 1485

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1490                 1495                 1500

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1505                 1510                 1515
```

```
Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1520                1525                1530

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1535                1540                1545

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1550                1555                1560

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1565                1570                1575

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1580                1585                1590

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1595                1600                1605

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1610                1615                1620

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1625                1630                1635

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1640                1645                1650

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1655                1660                1665

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1670                1675                1680

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1685                1690                1695

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1700                1705                1710

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1715                1720                1725

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1730                1735                1740

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1745                1750                1755

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1760                1765                1770
```

Arg Ser Asn Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp
1775 1780 1785

Leu Gln Val Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr
1790 1795 1800

Thr Gln Gly Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu
1805 1810 1815

Phe Leu Ile Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe
1820 1825 1830

Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser
1835 1840 1845

Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg
1850 1855 1860

Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu
1865 1870 1875

Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala
1880 1885 1890

Gly Ser Pro Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu
1895 1900 1905

Glu Asp Leu Ser Pro Ala Thr Gly His His His His His His
1910 1915 1920

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<211> 5769
<212> DNA
<213> Artificial Sequence

<220>
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ccaggttcgt caccctcagc atcaacgggg acgggaccgg gggcgtcacc aggaacgtcc 180

tccaccggct cgccgggtgc atcacccgga acgtcatcga ccggatcgcc agggagctcg 240

acgccatcag gcgcaacagg atcacctggc tcaagcccta gcgcgtcaac cggcacgggt 300

ccgggtgcct cccctggcac gtccagcacc ggatcacccg gatcgagccc atccgcctca 360

accggaaccg gacccggtac accagggtcg ggaacagcct cctcgtcacc aggctcctca 420

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<211> 1919
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Gly Ser Ser
20 25 30

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
35 40 45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
50 55 60

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Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65                  70                  75                  80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
                85                  90                  95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
            100                 105                 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
        115                 120                 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
    130                 135                 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145                 150                 155                 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335
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Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605
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Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880
```

```
Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
        915                 920                 925

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
    930                 935                 940

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
945                 950                 955                 960

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
                965                 970                 975

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly
            980                 985                 990

Thr Ser Ser Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
        995                 1000                1005

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1010                1015                1020

Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser
    1025                1030                1035

Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    1040                1045                1050

Ser Ser  Thr Gly Ser Pro Ala  Ser Ser Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140
```

```
Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1295                1300                1305

Pro Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    1310                1315                1320

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1325                1330                1335

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1340                1345                1350

Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1355                1360                1365

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1370                1375                1380

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Pro  Ala Gly Ser
    1385                1390                1395
```

```
Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1400                1405                1410

Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1415                1420                1425

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1430                1435                1440

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ala Ser  Ser Ala Pro
    1445                1450                1455

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                1555                1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                1615                1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650
```

```
Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1880                1885                1890

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
    1895                1900                1905
```

Ser Pro Ala Thr Gly His His His His His His
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<212> DNA
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tccaccggct cgccgggtgc atcacccgga acgtcatcga ccggatcgcc agggagctcg 240

acgccatcag gcgcaacagg atcacctggc tcaagcccta gcgcgtcaac cggcacgggt 300

ccgggtgcct cccctggcac gtccagcacc ggatcacccg gatcgagccc atccgcctca 360

accggaaccg gacccggtac accagggtcg ggaacagcct cctcgtcacc aggctcctca 420

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<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic polypeptide

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Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
35 40 45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
50 55 60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65 70 75 80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
85 90 95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
100 105 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
115 120 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
130 135 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145 150 155 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

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Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495
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Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765
```

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Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
            980                 985                 990

Ser Pro Gly Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
        995                 1000                1005

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1010                1015                1020

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1025                1030                1035
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Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
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Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
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Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1070                1075                1080

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1085                1090                1095

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1100                1105                1110

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
    1115                1120                1125

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1130                1135                1140

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1145                1150                1155

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1160                1165                1170

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1175                1180                1185

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1190                1195                1200

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1205                1210                1215

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1220                1225                1230

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1235                1240                1245

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1250                1255                1260

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1265                1270                1275

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1280                1285                1290
```

```
Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1295                1300                1305

Arg Gly  Ala Pro Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1310                1315                1320

Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser
    1325                1330                1335

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
    1340                1345                1350

Ser Gly  Ser Glu Thr Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1355                1360                1365

Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1370                1375                1380

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Pro
    1385                1390                1395

Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Ser Pro  Ala Gly Ser
    1400                1405                1410

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1415                1420                1425

Thr Glu  Glu Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1430                1435                1440

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Ala Ser
    1445                1450                1455

Ser Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1460                1465                1470

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1475                1480                1485

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1490                1495                1500

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1505                1510                1515

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1520                1525                1530

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1535                1540                1545
```

```
Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1550                1555                1560

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1565                1570                1575

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1580                1585                1590

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1595                1600                1605

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1610                1615                1620

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1625                1630                1635

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1640                1645                1650

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1655                1660                1665

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1670                1675                1680

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1685                1690                1695

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1700                1705                1710

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1715                1720                1725

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1730                1735                1740

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1745                1750                1755

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1760                1765                1770

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1775                1780                1785

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1790                1795                1800
```

Thr Gln Gly Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu
1805 1810 1815

Phe Leu Ile Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe
1820 1825 1830

Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser
1835 1840 1845

Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg
1850 1855 1860

Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu
1865 1870 1875

Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala
1880 1885 1890

Gly Ser Pro Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu
1895 1900 1905

Glu Asp Leu Ser Pro Ala Thr Gly His His His His His His
1910 1915 1920

<210> 1046
<211> 5769
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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ccaggttcgt caccctcagc atcaacgggg acgggaccgg gggcgtcacc aggaacgtcc 180

tccaccggct cgccgggtgc atcacccgga acgtcatcga ccggatcgcc agggagctcg 240

acgccatcag gcgcaacagg atcacctggc tcaagcccta gcgcgtcaac cggcacgggt 300

ccgggtgcct cccctggcac gtccagcacc ggatcacccg gatcgagccc atccgcctca 360

accggaaccg gacccggtac accagggtcg ggaacagcct cctcgtcacc aggctcctca 420

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cccggagcat cgccaggaac ttcgagcacg gggtcgcccg cctcgagcga cgcaagattt 540

cctcctagag tgccaaaatc ttttccattc aacacctcag tcgtgtacaa aaagactctg 600

tttgtagaat tcacggatca ccttttcaac atcgctaagc caaggccacc ctggatgggt 660

ctgctaggtc ctaccatcca ggctgaggtt tatgatacag tggtcattac acttaagaac 720

atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag 780

ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct 840

ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

ggcctcattg gagccctact agtatgtaga gaagggagtc tggccaagga aaagacacag 1020

accttgcaca aatttatact acttttgct gtatttgatg aagggaaaag ttggcactca 1080

gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

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ataactttcc ttactgctca aacactcttg atggaccttg gacagtttct actgttttgt 1380

catatctctt cccaccaaca tgatggcatg gaagcttatg tcaaagtaga cagctgtcca 1440

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cctttacttt atggggaagt tggagacaca ctgttgatta tatttaagaa tcaagcaagc 1860

agaccatata acatctaccc tcacggaatc actgatgtcc gtcctttgta ttcaaggaga 1920

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gccgaacaaa aactcatctc agaagaggat ctgtcacctg caaccggtca tcatcaccat 5760

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<210> 1047
<211> 1922
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

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Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365
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```
Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
                405                 410                 415

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            420                 425                 430

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
        435                 440                 445

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    450                 455                 460

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
465                 470                 475                 480

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
                485                 490                 495

Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            500                 505                 510

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
        515                 520                 525

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
    530                 535                 540

Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640
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```
Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910
```

```
Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
        915                 920                 925

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
    930                 935                 940

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
945                 950                 955                 960

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
                965                 970                 975

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly
            980                 985                 990

Thr Ser Ser Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
        995                 1000                1005

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1010                1015                1020

Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser
    1025                1030                1035

Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    1040                1045                1050

Ser Ser  Thr Gly Ser Pro Ala  Ser Ser Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Gly  Ala Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1130                1135                1140

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1145                1150                1155

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1160                1165                1170
```

```
Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1175                1180                1185

Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1190                1195                1200

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1205                1210                1215

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1220                1225                1230

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1235                1240                1245

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1250                1255                1260

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
    1265                1270                1275

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1280                1285                1290

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1295                1300                1305

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1310                1315                1320

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1325                1330                1335

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1340                1345                1350

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1355                1360                1365

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1370                1375                1380

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1385                1390                1395

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1400                1405                1410

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1415                1420                1425
```

```
Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1430                1435                1440

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1445                1450                1455

Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1460                1465                1470

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1475                1480                1485

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1490                1495                1500

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1505                1510                1515

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1520                1525                1530

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1535                1540                1545

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1550                1555                1560

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1565                1570                1575

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1580                1585                1590

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1595                1600                1605

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1610                1615                1620

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1625                1630                1635

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1640                1645                1650

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1655                1660                1665

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1670                1675                1680
```

```
Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1685                1690                1695

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1700                1705                1710

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1715                1720                1725

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1730                1735                1740

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1745                1750                1755

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1760                1765                1770

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1775                1780                1785

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1790                1795                1800

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1805                1810                1815

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1820                1825                1830

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1835                1840                1845

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    1850                1855                1860

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    1865                1870                1875

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    1880                1885                1890

Gly Ser  Pro Gly Ala Glu Thr  Ala Glu Gln Lys Leu  Ile Ser Glu
    1895                1900                1905

Glu Asp  Leu Ser Pro Ala Thr  Gly His His His His  His His
    1910                1915                1920
```

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<211> 5769
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

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atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

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gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

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gcccccgatg gcgcgccaac cagtacggag ccgtccgagg ggagcgcacc aggaagcccg 1260

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gaatcggcaa ccccagaatc agggccagga tctgagcccg cgacttcggg ctccgagacg 1500

cctgggacat ccaccgagcc ctccgaagga tcagccccag gcaccagcac ggagccctct 1560

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tatttgaaca atggccctca gcggattggt aggaagtaca aaaaagtccg atttatggca 1740

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<210> 1049
<211> 1919
<212> PRT

<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His

```
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
                405                 410                 415

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            420                 425                 430

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
        435                 440                 445

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    450                 455                 460

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
465                 470                 475                 480

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
                485                 490                 495

Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            500                 505                 510

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
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515 520 525

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
530 535 540

Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545 550 555 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
565 570 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
580 585 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
595 600 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
610 615 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
645 650 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser

785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
835 840 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
885 890 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
900 905 910

Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
915 920 925

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
930 935 940

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
945 950 955 960

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
965 970 975

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly
980 985 990

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
995 1000 1005

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
1010 1015 1020

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
1025 1030 1035

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
1040 1045 1050

Ser Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Gln Glu Glu Ile

1055 1060 1065

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
1070 1075 1080

Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln
1085 1090 1095

Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp
1100 1105 1110

Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala
1115 1120 1125

Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu
1130 1135 1140

Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
1145 1150 1155

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1160 1165 1170

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1175 1180 1185

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1190 1195 1200

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1205 1210 1215

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1220 1225 1230

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1235 1240 1245

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1250 1255 1260

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1265 1270 1275

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1280 1285 1290

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Gly Ala
1295 1300 1305

Pro Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu

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Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1325                1330                1335

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1340                1345                1350

Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1355                1360                1365

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1370                1375                1380

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Pro  Ala Gly Ser
    1385                1390                1395

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1400                1405                1410

Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1415                1420                1425

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1430                1435                1440

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ala Ser  Ser Ala Pro
    1445                1450                1455

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                1555                1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
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Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1580 1585 1590

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1595 1600 1605

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1610 1615 1620

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1625 1630 1635

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1640 1645 1650

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1655 1660 1665

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1670 1675 1680

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1685 1690 1695

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1700 1705 1710

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1715 1720 1725

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1730 1735 1740

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1745 1750 1755

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1760 1765 1770

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1775 1780 1785

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1790 1795 1800

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1805 1810 1815

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn

1820 1825 1830

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1835 1840 1845

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1850 1855 1860

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1865 1870 1875

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1880 1885 1890

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1895 1900 1905

Ser Pro Ala Thr Gly His His His His His His
1910 1915

<210> 1050
<211> 5760
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<220>
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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

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ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

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tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

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atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

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cctgggacat ccaccgagcc ctccgaagga tcagccccag gcaccagcac ggagccctct 1560

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cctttacttt atggggaagt tggagacaca ctgttgatta tatttaagaa tcaagcaagc 1860

agaccatata acatctaccc tcacggaatc actgatgtcc gtcctttgta ttcaaggaga 1920

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<211> 1922
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

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Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400
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Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
               405                 410                 415

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            420                 425                 430

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
        435                 440                 445

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    450                 455                 460

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
465                 470                 475                 480

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
               485                 490                 495

Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            500                 505                 510

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
        515                 520                 525

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
    530                 535                 540

Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
               565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
               645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670
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Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940
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Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
            980                 985                 990

Ser Pro Gly Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
        995                 1000                1005

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1010                1015                1020

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1025                1030                1035

Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1040                1045                1050

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1055                1060                1065

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1070                1075                1080

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1085                1090                1095

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1100                1105                1110

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
    1115                1120                1125

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1130                1135                1140

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1145                1150                1155

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1160                1165                1170

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1175                1180                1185

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1190                1195                1200
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Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1205                 1210                 1215

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1220                 1225                 1230

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1235                 1240                 1245

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1250                 1255                 1260

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1265                 1270                 1275

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1280                 1285                 1290

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1295                 1300                 1305

Arg Gly  Ala Pro Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1310                 1315                 1320

Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser
    1325                 1330                 1335

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
    1340                 1345                 1350

Ser Gly  Ser Glu Thr Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1355                 1360                 1365

Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1370                 1375                 1380

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Pro
    1385                 1390                 1395

Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Ser Pro  Ala Gly Ser
    1400                 1405                 1410

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1415                 1420                 1425

Thr Glu  Glu Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1430                 1435                 1440

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Ala Ser
    1445                 1450                 1455
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Ser Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1460                1465                1470

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1475                1480                1485

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1490                1495                1500

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1505                1510                1515

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1520                1525                1530

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1535                1540                1545

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1550                1555                1560

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1565                1570                1575

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1580                1585                1590

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1595                1600                1605

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1610                1615                1620

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1625                1630                1635

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1640                1645                1650

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1655                1660                1665

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1670                1675                1680

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1685                1690                1695

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1700                1705                1710
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His Tyr Ser Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys
1715 1720 1725

Asp Leu Asn Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala
1730 1735 1740

Ile Ser Asp Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met
1745 1750 1755

Phe Ala Thr Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly
1760 1765 1770

Arg Ser Asn Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp
1775 1780 1785

Leu Gln Val Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr
1790 1795 1800

Thr Gln Gly Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu
1805 1810 1815

Phe Leu Ile Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe
1820 1825 1830

Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser
1835 1840 1845

Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg
1850 1855 1860

Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu
1865 1870 1875

Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala
1880 1885 1890

Gly Ser Pro Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu
1895 1900 1905

Glu Asp Leu Ser Pro Ala Thr Gly His His His His His His
1910 1915 1920

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<213> Artificial Sequence

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

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35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
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Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
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Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

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275 280 285

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Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
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Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
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Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
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Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
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Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

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Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
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His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp

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Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
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Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
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Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
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Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
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Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
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Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
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Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Gly Ala Pro Gly Thr Pro
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Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser
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Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
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Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
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Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr
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Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
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Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
850 855 860

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly
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Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
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915 920 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
930 935 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
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Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
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980 985 990

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Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
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Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
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Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
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Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1175                1180                1185

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
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Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
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Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
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Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
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Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
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Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
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Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
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Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
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Arg Gly  Ala Pro Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
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Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser
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Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
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Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
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Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
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Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
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Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
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His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
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Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
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Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
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Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
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Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
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Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
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His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
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Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
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Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu
1865 1870 1875

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Gly Ser Ser
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Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
35 40 45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
50 55 60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65 70 75 80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
85 90 95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
100 105 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
115 120 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
130 135 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145 150 155 160

```
Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430
```

```
Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
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Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu
545                 550                 555                 560

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
                565                 570                 575

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            580                 585                 590

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
        595                 600                 605

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    610                 615                 620

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
625                 630                 635                 640

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
                645                 650                 655

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
            660                 665                 670

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
        675                 680                 685

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp
    690                 695                 700
```

```
Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975
```

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
980 985 990

His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val
995 1000 1005

Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr
1010 1015 1020

Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu
1025 1030 1035

Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His Gln
1040 1045 1050

Arg Glu Ile Thr Arg Thr Thr Leu Gln Gly Ala Pro Gly Thr Pro
1055 1060 1065

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr
1070 1075 1080

Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
1085 1090 1095

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
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Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
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Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr
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Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
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Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
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Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
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Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
1190 1195 1200

Ser Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Gln Glu Glu Ile
1205 1210 1215

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
1220 1225 1230

```
Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1235                 1240                 1245

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1250                 1255                 1260

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1265                 1270                 1275

Gln Gly  Ala Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1280                 1285                 1290

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1295                 1300                 1305

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1310                 1315                 1320

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1325                 1330                 1335

Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1340                 1345                 1350

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1355                 1360                 1365

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1370                 1375                 1380

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1385                 1390                 1395

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1400                 1405                 1410

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
    1415                 1420                 1425

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1430                 1435                 1440

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1445                 1450                 1455

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1460                 1465                 1470

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1475                 1480                 1485
```

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Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
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Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1505                1510                1515

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1520                1525                1530

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1535                1540                1545

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1550                1555                1560

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1565                1570                1575

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1580                1585                1590

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1595                1600                1605

Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1610                1615                1620

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1625                1630                1635

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1640                1645                1650

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1655                1660                1665

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1670                1675                1680

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1685                1690                1695

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1700                1705                1710

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1715                1720                1725

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1730                1735                1740
```

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Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1745                 1750                 1755

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1760                 1765                 1770

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1775                 1780                 1785

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1790                 1795                 1800

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1805                 1810                 1815

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1820                 1825                 1830

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1835                 1840                 1845

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1850                 1855                 1860

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1865                 1870                 1875

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1880                 1885                 1890

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1895                 1900                 1905

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1910                 1915                 1920

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1925                 1930                 1935

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1940                 1945                 1950

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1955                 1960                 1965

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1970                 1975                 1980

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1985                 1990                 1995
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Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg
2000 2005 2010

Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu
2015 2020 2025

Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala
2030 2035 2040

Gly Ser Pro Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu
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Glu Asp Leu Ser Pro Ala Thr Gly His His His His His His
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<213> Artificial Sequence

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Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
35 40 45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
50 55 60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65 70 75 80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser

85 90 95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
100 105 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
115 120 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
130 135 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145 150 155 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
260 265 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
275 280 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
290 295 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305 310 315 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
325 330 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
340 345 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln

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Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu
545                 550                 555                 560

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
                565                 570                 575

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            580                 585                 590

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
        595                 600                 605

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    610                 615                 620

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
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625 630 635 640

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
645 650 655

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
660 665 670

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
675 680 685

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp
690 695 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705 710 715 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
725 730 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
740 745 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
755 760 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
770 775 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785 790 795 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
805 810 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
820 825 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
835 840 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
850 855 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865 870 875 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
885 890 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn

900 905 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
915 920 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
930 935 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945 950 955 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
965 970 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
980 985 990

His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val
995 1000 1005

Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr
1010 1015 1020

Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu
1025 1030 1035

Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His Gln
1040 1045 1050

Arg Glu Ile Thr Arg Thr Thr Leu Gln Gly Ala Pro Gly Thr Pro
1055 1060 1065

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr
1070 1075 1080

Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
1085 1090 1095

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
1100 1105 1110

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
1115 1120 1125

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1145 1150 1155

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro

1160 1165 1170

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
1175 1180 1185

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
1190 1195 1200

Ser Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Gln Glu Glu Ile
1205 1210 1215

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
1220 1225 1230

Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln
1235 1240 1245

Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp
1250 1255 1260

Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala
1265 1270 1275

Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu
1280 1285 1290

Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
1295 1300 1305

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1310 1315 1320

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1325 1330 1335

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1340 1345 1350

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1355 1360 1365

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1370 1375 1380

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1385 1390 1395

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1400 1405 1410

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr

1415 1420 1425

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1430 1435 1440

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Gly Ala
1445 1450 1455

Pro Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
1460 1465 1470

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala
1475 1480 1485

Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
1490 1495 1500

Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
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Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
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Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser
1550 1555 1560

Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
1565 1570 1575

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Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Ala Pro
1595 1600 1605

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1610 1615 1620

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1625 1630 1635

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1640 1645 1650

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1655 1660 1665

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn

1670 1675 1680

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1685 1690 1695

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1700 1705 1710

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1715 1720 1725

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1730 1735 1740

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1745 1750 1755

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1760 1765 1770

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1775 1780 1785

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1790 1795 1800

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1805 1810 1815

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1820 1825 1830

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1835 1840 1845

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1850 1855 1860

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1865 1870 1875

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1880 1885 1890

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1895 1900 1905

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1910 1915 1920

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val

1925 1930 1935

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1940 1945 1950

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1955 1960 1965

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1970 1975 1980

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1985 1990 1995

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
2000 2005 2010

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
2015 2020 2025

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
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Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
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Ser Pro Ala Thr Gly His His His His His His
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Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
        35                  40                  45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
    50                  55                  60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65                  70                  75                  80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
                85                  90                  95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
            100                 105                 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
        115                 120                 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
    130                 135                 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145                 150                 155                 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285
```

```
Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu
545                 550                 555                 560
```

```
Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
               565                 570                 575

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            580                 585                 590

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
        595                 600                 605

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    610                 615                 620

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
625                 630                 635                 640

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
               645                 650                 655

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
            660                 665                 670

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
        675                 680                 685

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp
    690                 695                 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
               725                 730                 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
               805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830
```

```
Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020

Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095
```

```
Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Gly  Ala Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1130                1135                1140

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1145                1150                1155

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1160                1165                1170

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1175                1180                1185

Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1190                1195                1200

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1205                1210                1215

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1220                1225                1230

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1235                1240                1245

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1250                1255                1260

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
    1265                1270                1275

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1280                1285                1290

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1295                1300                1305

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1310                1315                1320

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1325                1330                1335

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1340                1345                1350
```

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Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
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Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1370                1375                1380

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1385                1390                1395

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1400                1405                1410

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1415                1420                1425

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1430                1435                1440

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1445                1450                1455

Arg Gly  Ala Pro Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1460                1465                1470

Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser
    1475                1480                1485

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
    1490                1495                1500

Ser Gly  Ser Glu Thr Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1505                1510                1515

Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1520                1525                1530

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Pro
    1535                1540                1545

Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Ser Pro  Ala Gly Ser
    1550                1555                1560

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1565                1570                1575

Thr Glu  Glu Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1580                1585                1590

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Ala Ser
    1595                1600                1605
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Ser Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
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Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
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Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1640                1645                1650

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
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Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1670                1675                1680

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1685                1690                1695

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1700                1705                1710

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1715                1720                1725

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1730                1735                1740

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1745                1750                1755

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1760                1765                1770

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1775                1780                1785

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1790                1795                1800

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1805                1810                1815

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1820                1825                1830

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1835                1840                1845

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1850                1855                1860
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His Tyr Ser Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys
1865 1870 1875

Asp Leu Asn Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala
1880 1885 1890

Ile Ser Asp Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met
1895 1900 1905

Phe Ala Thr Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly
1910 1915 1920

Arg Ser Asn Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp
1925 1930 1935

Leu Gln Val Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr
1940 1945 1950

Thr Gln Gly Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu
1955 1960 1965

Phe Leu Ile Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe
1970 1975 1980

Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser
1985 1990 1995

Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg
2000 2005 2010

Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu
2015 2020 2025

Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala
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Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
35 40 45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
50 55 60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65 70 75 80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
85 90 95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
100 105 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
115 120 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
130 135 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145 150 155 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

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Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
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Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495
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Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
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Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765
```

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Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
        915                 920                 925

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
    930                 935                 940

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
945                 950                 955                 960

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
                965                 970                 975

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly
            980                 985                 990

Thr Ser Ser Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
        995                 1000                1005

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
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Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser
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Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
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Ser Ser  Thr Gly Ser Pro Ala  Ser Ser Ser Asp Gln  Glu Glu Ile
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Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Gly  Ala Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1130                1135                1140

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1145                1150                1155

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1160                1165                1170

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1175                1180                1185

Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1190                1195                1200

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1205                1210                1215

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1220                1225                1230

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1235                1240                1245

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
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Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
    1265                1270                1275

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1280                1285                1290
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Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
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Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1310                1315                1320

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1325                1330                1335

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1340                1345                1350

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1355                1360                1365

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1370                1375                1380

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1385                1390                1395

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1400                1405                1410

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1415                1420                1425

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1430                1435                1440

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1445                1450                1455

Arg Gly  Ala Pro Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1460                1465                1470

Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser
    1475                1480                1485

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
    1490                1495                1500

Ser Gly  Ser Glu Thr Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1505                1510                1515

Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1520                1525                1530

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Pro
    1535                1540                1545
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Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Ser Pro  Ala Gly Ser
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Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1565                1570                1575

Thr Glu  Glu Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1580                1585                1590

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Ala Ser
    1595                1600                1605

Ser Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1610                1615                1620

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1625                1630                1635

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1640                1645                1650

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1655                1660                1665

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1670                1675                1680

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1685                1690                1695

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1700                1705                1710

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1715                1720                1725

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1730                1735                1740

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1745                1750                1755

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1760                1765                1770

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1775                1780                1785

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1790                1795                1800
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Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1805                1810                1815

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1820                1825                1830

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1835                1840                1845

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1850                1855                1860

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1865                1870                1875

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1880                1885                1890

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1895                1900                1905

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1910                1915                1920

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1925                1930                1935

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1940                1945                1950

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1955                1960                1965

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1970                1975                1980

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1985                1990                1995

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    2000                2005                2010

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    2015                2020                2025

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    2030                2035                2040

Gly Ser  Pro Gly Ala Glu Thr  Ala Glu Gln Lys Leu  Ile Ser Glu
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Glu Asp Leu Ser Pro Ala Thr Gly His His His His His His
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

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His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
                405                 410                 415

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            420                 425                 430
```

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Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
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Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    450                 455                 460

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
465                 470                 475                 480

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
                485                 490                 495

Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            500                 505                 510

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
        515                 520                 525

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
    530                 535                 540

Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700
```

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Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
        915                 920                 925

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
    930                 935                 940

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
945                 950                 955                 960

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
                965                 970                 975
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Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly
           980                 985                 990

Thr Ser Ser Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
        995                 1000                1005

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1010                1015                1020

Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser
    1025                1030                1035

Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    1040                1045                1050

Ser Ser  Thr Gly Ser Pro Ala  Ser Ser Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Gly  Ala Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1130                1135                1140

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1145                1150                1155

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1160                1165                1170

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1175                1180                1185

Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1190                1195                1200

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1205                1210                1215

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1220                1225                1230
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Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
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Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1250                 1255                 1260

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
    1265                 1270                 1275

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1280                 1285                 1290

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1295                 1300                 1305

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1310                 1315                 1320

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1325                 1330                 1335

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1340                 1345                 1350

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1355                 1360                 1365

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1370                 1375                 1380

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1385                 1390                 1395

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1400                 1405                 1410

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1415                 1420                 1425

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1430                 1435                 1440

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1445                 1450                 1455

Arg Gly  Ala Pro Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1460                 1465                 1470

Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser
    1475                 1480                 1485
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Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
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Ser Gly  Ser Glu Thr Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1505                 1510                 1515

Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1520                 1525                 1530

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Pro
    1535                 1540                 1545

Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Ser Pro  Ala Gly Ser
    1550                 1555                 1560

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1565                 1570                 1575

Thr Glu  Glu Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1580                 1585                 1590

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Ala Ser
    1595                 1600                 1605

Ser Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1610                 1615                 1620

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1625                 1630                 1635

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1640                 1645                 1650

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1655                 1660                 1665

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1670                 1675                 1680

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1685                 1690                 1695

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1700                 1705                 1710

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
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Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1730                 1735                 1740
```

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Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1745                 1750                 1755

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1760                 1765                 1770

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1775                 1780                 1785

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1790                 1795                 1800

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1805                 1810                 1815

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1820                 1825                 1830

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1835                 1840                 1845

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1850                 1855                 1860

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1865                 1870                 1875

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1880                 1885                 1890

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1895                 1900                 1905

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1910                 1915                 1920

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1925                 1930                 1935

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1940                 1945                 1950

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1955                 1960                 1965

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1970                 1975                 1980

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1985                 1990                 1995
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Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg
2000 2005 2010

Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu
2015 2020 2025

Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala
2030 2035 2040

Gly Ser Pro Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu
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Glu Asp Leu Ser Pro Ala Thr Gly His His His His His His
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35 40 45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
50 55 60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65 70 75 80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser

85 90 95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
100 105 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
115 120 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
130 135 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145 150 155 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
260 265 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
275 280 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
290 295 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305 310 315 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
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Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
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Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln

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Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
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Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
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Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
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Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu
545                 550                 555                 560

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
                565                 570                 575

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            580                 585                 590

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
        595                 600                 605

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    610                 615                 620

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
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625 630 635 640

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645 650 655

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
660 665 670

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
675 680 685

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp
690 695 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705 710 715 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
725 730 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
740 745 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
755 760 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
770 775 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785 790 795 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
805 810 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
820 825 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
835 840 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
850 855 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865 870 875 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
885 890 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn

900 905 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
915 920 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
930 935 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945 950 955 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
965 970 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
980 985 990

His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val
995 1000 1005

Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr
1010 1015 1020

Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu
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Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His Gln
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Arg Glu Ile Thr Arg Thr Thr Leu Gln Gly Ala Pro Gly Thr Pro
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1160 1165 1170

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Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
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Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln
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Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp
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Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala
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Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
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Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
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Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
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Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
1400 1405 1410

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser

1415 1420 1425

Ser Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val
1430 1435 1440

Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg
1445 1450 1455

Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
1460 1465 1470

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala
1475 1480 1485

Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu
1490 1495 1500

Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro
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Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala
1520 1525 1530

Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser
1535 1540 1545

Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
1550 1555 1560

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg
1565 1570 1575

Gln Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp
1580 1585 1590

Glu Thr Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys
1595 1600 1605

Arg Gly Ala Pro Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
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Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser
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1640 1645 1650

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1670 1675 1680

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro
1685 1690 1695

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser
1700 1705 1710

Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser
1715 1720 1725

Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
1730 1735 1740

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser
1745 1750 1755

Ser Ala Pro Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu
1760 1765 1770

Asn Tyr Arg Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu
1775 1780 1785

Pro Gly Leu Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu
1790 1795 1800

Leu Ser Met Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser
1805 1810 1815

Gly His Val Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala
1820 1825 1830

Leu Tyr Asn Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu
1835 1840 1845

Pro Ser Lys Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu
1850 1855 1860

His Leu His Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn
1865 1870 1875

Lys Cys Gln Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp
1880 1885 1890

Phe Gln Ile Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys
1895 1900 1905

Leu Ala Arg Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr
1910 1915 1920

Lys Glu Pro Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met

1925 1930 1935

Ile Ile His Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser
1940 1945 1950

Ser Leu Tyr Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly
1955 1960 1965

Lys Lys Trp Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met
1970 1975 1980

Val Phe Phe Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile
1985 1990 1995

Phe Asn Pro Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr
2000 2005 2010

His Tyr Ser Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys
2015 2020 2025

Asp Leu Asn Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala
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Ile Ser Asp Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met
2045 2050 2055

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2060 2065 2070

Arg Ser Asn Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp
2075 2080 2085

Leu Gln Val Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr
2090 2095 2100

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2105 2110 2115

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Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg
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Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu
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Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala

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<211> 2053
<212> PRT
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Met Gln Gly Ala Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
35 40 45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
50 55 60

```
Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65                  70                  75                  80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
                85                  90                  95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
            100                 105                 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
        115                 120                 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
    130                 135                 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145                 150                 155                 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335
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Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605
```

```
Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880
```

```
Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140
```

```
Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395
```

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Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
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Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650
```

```
Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                 1660                 1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                 1675                 1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                 1690                 1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                 1705                 1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                 1720                 1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1730                 1735                 1740

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1745                 1750                 1755

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1760                 1765                 1770

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1775                 1780                 1785

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1790                 1795                 1800

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1805                 1810                 1815

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1820                 1825                 1830

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1835                 1840                 1845

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1850                 1855                 1860

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    1865                 1870                 1875

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    1880                 1885                 1890

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1895                 1900                 1905
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Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1910 1915 1920

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1925 1930 1935

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
1940 1945 1950

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
1955 1960 1965

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
1970 1975 1980

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
1985 1990 1995

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2000 2005 2010

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2015 2020 2025

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Ala Thr Gly His His His His His His
2045 2050

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<211> 2053
<212> PRT
<213> Artificial Sequence

<220>
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Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
35 40 45

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
50 55 60

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
65 70 75 80

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
85 90 95

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
100 105 110

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
115 120 125

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
130 135 140

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
145 150 155 160

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ala Ser Ser
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
260 265 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
275 280 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu

290 295 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305 310 315 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
325 330 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
340 345 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
355 360 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
370 375 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385 390 395 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
405 410 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
420 425 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
435 440 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
450 455 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465 470 475 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
485 490 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
500 505 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
515 520 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
530 535 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545 550 555 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val

565 570 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
580 585 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
595 600 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
610 615 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
645 650 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg

835 840 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
885 890 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
900 905 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
915 920 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
930 935 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945 950 955 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
965 970 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
980 985 990

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
995 1000 1005

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser
1010 1015 1020

Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
1025 1030 1035

Ser Ala Pro Gly Ala Ser Ser Pro Pro Val Leu Lys Arg His Gln
1040 1045 1050

Ala Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
1055 1060 1065

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
1070 1075 1080

Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln
1085 1090 1095

Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp

1100 1105 1110

Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala
1115 1120 1125

Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu
1130 1135 1140

Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
1145 1150 1155

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1160 1165 1170

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1175 1180 1185

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1190 1195 1200

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1205 1210 1215

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1220 1225 1230

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1235 1240 1245

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1250 1255 1260

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1265 1270 1275

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1280 1285 1290

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1295 1300 1305

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1310 1315 1320

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1325 1330 1335

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1340 1345 1350

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val

1355 1360 1365

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1370 1375 1380

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1385 1390 1395

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1400 1405 1410

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1415 1420 1425

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1430 1435 1440

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1445 1450 1455

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1460 1465 1470

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1475 1480 1485

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1490 1495 1500

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1505 1510 1515

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1520 1525 1530

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1535 1540 1545

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1550 1555 1560

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1565 1570 1575

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1580 1585 1590

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1595 1600 1605

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn

1610 1615 1620

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1625 1630 1635

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Pro Gly Thr
1730 1735 1740

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1745 1750 1755

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
1760 1765 1770

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1775 1780 1785

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1790 1795 1800

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
1805 1810 1815

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
1820 1825 1830

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1835 1840 1845

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
1850 1855 1860

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly

1865 1870 1875

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
1880 1885 1890

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1895 1900 1905

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1910 1915 1920

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1925 1930 1935

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
1940 1945 1950

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
1955 1960 1965

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
1970 1975 1980

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
1985 1990 1995

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2000 2005 2010

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2015 2020 2025

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Ala Thr Gly His His His His His His
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<210> 1071
<211> 2053
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Gly Ser Ser
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Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
35 40 45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
50 55 60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65 70 75 80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
85 90 95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
100 105 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
115 120 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
130 135 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145 150 155 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

```
Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525
```

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Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800
```

```
Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065
```

```
Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320
```

```
Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575
```

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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                 1585                 1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                 1600                 1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                 1615                 1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                 1630                 1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                 1645                 1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                 1660                 1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                 1675                 1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                 1690                 1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                 1705                 1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                 1720                 1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1730                 1735                 1740

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1745                 1750                 1755

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1760                 1765                 1770

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1775                 1780                 1785

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1790                 1795                 1800

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1805                 1810                 1815

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1820                 1825                 1830
```

```
Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1850                 1855                 1860

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    1865                 1870                 1875

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    1880                 1885                 1890

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1895                 1900                 1905

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1910                 1915                 1920

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1925                 1930                 1935

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1940                 1945                 1950

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    1955                 1960                 1965

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    1970                 1975                 1980

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Glu Pro Ala
    1985                 1990                 1995

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2000                 2005                 2010

Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    2015                 2020                 2025

Pro Ala  Ser Ser Glu Gln Lys  Leu Ile Ser Glu Glu  Asp Leu Ser
    2030                 2035                 2040

Pro Ala  Thr Gly His His His  His His His
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<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic polynucleotide

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<210> 1073
<211> 2053
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1073

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Gly Ala Pro Gly Ser Ser Pro Ser
35 40 45

Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
50 55 60

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65 70 75 80

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro
85 90 95

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
100 105 110

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
115 120 125

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser
130 135 140

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145 150 155 160

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
165 170 175

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

```
Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480
```

```
Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750
```

```
Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020
```

```
Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
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Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275
```

```
Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530
```

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Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1730                1735                1740

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1745                1750                1755

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1760                1765                1770

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1775                1780                1785
```

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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1790                 1795                 1800

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1805                 1810                 1815

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1820                 1825                 1830

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1835                 1840                 1845

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1850                 1855                 1860

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    1865                 1870                 1875

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    1880                 1885                 1890

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1895                 1900                 1905

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1910                 1915                 1920

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1925                 1930                 1935

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1940                 1945                 1950

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    1955                 1960                 1965

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    1970                 1975                 1980

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Glu Pro Ala
    1985                 1990                 1995

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2000                 2005                 2010

Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    2015                 2020                 2025

Pro Ala  Ser Ser Glu Gln Lys  Leu Ile Ser Glu Glu  Asp Leu Ser
    2030                 2035                 2040
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Pro Ala Thr Gly His His His His His His
2045 2050

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<211> 2053
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Gly Ala Pro Thr Ser Glu Ser Ala
35 40 45

Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu
50 55 60

Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
65 70 75 80

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala
85 90 95

Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
100 105 110

Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr
115 120 125

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr
130 135 140

Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
145 150 155 160

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Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser
                165                 170                 175

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ala Ser Ser Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430
```

```
Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700
```

```
Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975
```

```
Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230
```

```
Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485
```

```
Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1730                1735                1740
```

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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1745                1750                1755

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1760                1765                1770

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1775                1780                1785

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1790                1795                1800

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1805                1810                1815

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1820                1825                1830

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1835                1840                1845

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1850                1855                1860

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    1865                1870                1875

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    1880                1885                1890

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1895                1900                1905

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1910                1915                1920

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1925                1930                1935

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1940                1945                1950

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    1955                1960                1965

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    1970                1975                1980

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Glu Pro Ala
    1985                1990                1995
```

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2000 2005 2010

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2015 2020 2025

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
2030 2035 2040

Pro Ala Thr Gly His His His His His His
2045 2050

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ccaggaacat ctacagagcc ctctgaaggc tccgctccag ggtccccagc cggcagtccc 5460

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gagtcggcaa ccccggagtc tggacccgga acttccgaaa gtgccacacc agagtccggt 5760

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<210> 1077
<211> 2053
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1077

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20 25 30

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
35 40 45

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
50 55 60

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
65 70 75 80

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
85 90 95

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
100 105 110

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro

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        115                 120                 125

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
    130                 135                 140

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
145                 150                 155                 160

Ser Thr Glu Glu Gly Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
```

385 390 395 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
405 410 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
420 425 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
435 440 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
450 455 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465 470 475 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
485 490 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
500 505 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
515 520 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
530 535 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545 550 555 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
565 570 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
580 585 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
595 600 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
610 615 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
645 650 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr

660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
835 840 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
885 890 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
900 905 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
915 920 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly

930 935 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945 950 955 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
965 970 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
980 985 990

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
995 1000 1005

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser
1010 1015 1020

Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
1025 1030 1035

Ser Ala Pro Gly Ala Ser Ser Pro Pro Val Leu Lys Arg His Gln
1040 1045 1050

Ala Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
1055 1060 1065

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
1070 1075 1080

Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln
1085 1090 1095

Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp
1100 1105 1110

Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala
1115 1120 1125

Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu
1130 1135 1140

Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
1145 1150 1155

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1160 1165 1170

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1175 1180 1185

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg

1190 1195 1200

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1205 1210 1215

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1220 1225 1230

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1235 1240 1245

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1250 1255 1260

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1265 1270 1275

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1280 1285 1290

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1295 1300 1305

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1310 1315 1320

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1325 1330 1335

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1340 1345 1350

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1355 1360 1365

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1370 1375 1380

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1385 1390 1395

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1400 1405 1410

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1415 1420 1425

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1430 1435 1440

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg

1445 1450 1455

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1460 1465 1470

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1475 1480 1485

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1490 1495 1500

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1505 1510 1515

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1520 1525 1530

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1535 1540 1545

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1550 1555 1560

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1565 1570 1575

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1580 1585 1590

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1595 1600 1605

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1610 1615 1620

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1625 1630 1635

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg

1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Pro Gly Thr
1730 1735 1740

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1745 1750 1755

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
1760 1765 1770

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1775 1780 1785

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1790 1795 1800

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
1805 1810 1815

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
1820 1825 1830

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1835 1840 1845

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
1850 1855 1860

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
1865 1870 1875

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
1880 1885 1890

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1895 1900 1905

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1910 1915 1920

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1925 1930 1935

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
1940 1945 1950

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu

1955 1960 1965

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
1970 1975 1980

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
1985 1990 1995

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2000 2005 2010

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2015 2020 2025

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Ala Thr Gly His His His His His His
2045 2050

<210> 1078
<211> 6162
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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tcggaaacgc caggcacaag tgagtctgca actcccgagt ccggacctgg ctccgagcct 180

gccactagcg gctccgagac tccgggaact tccgagagcg ctacaccaga aagcggaccc 240

ggaaccagta ccgaacctag cgagggctct gctccgggca gcccagccgg ctctcctaca 300

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gctacctcag gcagtgagac gccaggaacg agcgagtccg ctacaccgga gagtgggcca 420

gggagccctg ctggatctcc tacgtccact gaggaagggt caccagcggg ctcgcccacc 480

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<211> 2053
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Gly Ala Pro Gly Ser Ser Pro Ser
35 40 45

Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
50 55 60

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65 70 75 80

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Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro
                85                  90                  95

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
            100                 105                 110

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
        115                 120                 125

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser
    130                 135                 140

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145                 150                 155                 160

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
                165                 170                 175

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350
```

```
Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
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Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620
```

```
Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895
```

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Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
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Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155
```

```
Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
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Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410
```

```
Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665
```

```
Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1730                1735                1740

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1745                1750                1755

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1760                1765                1770

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1775                1780                1785

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1790                1795                1800

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1805                1810                1815

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1820                1825                1830

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1835                1840                1845

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1850                1855                1860

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    1865                1870                1875

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    1880                1885                1890

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1895                1900                1905

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1910                1915                1920
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Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1925 1930 1935

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
1940 1945 1950

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
1955 1960 1965

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
1970 1975 1980

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
1985 1990 1995

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2000 2005 2010

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Ala Thr Gly His His His His His His
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Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
    50                  55                  60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65                  70                  75                  80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
                85                  90                  95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
            100                 105                 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
        115                 120                 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
    130                 135                 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145                 150                 155                 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300
```

```
Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
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Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575
```

```
Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845
```

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Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110
```

```
Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
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Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365
```

```
Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620
```

```
Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1730                1735                1740

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1745                1750                1755

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1760                1765                1770

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1775                1780                1785

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1790                1795                1800

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1805                1810                1815

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1820                1825                1830

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1835                1840                1845

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1850                1855                1860

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    1865                1870                1875
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Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
1880 1885 1890

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1895 1900 1905

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1910 1915 1920

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1925 1930 1935

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
1940 1945 1950

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
1955 1960 1965

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
1970 1975 1980

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
1985 1990 1995

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2000 2005 2010

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Ala Thr Gly His His His His His His
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<212> DNA
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35 40 45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
50 55 60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
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Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
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Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
115 120 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
130 135 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145 150 155 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

```
Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525
```

```
Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800
```

```
Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065
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Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320
```

```
Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
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Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575
```

```
Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
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Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                 1600                 1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                 1615                 1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                 1630                 1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                 1645                 1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                 1660                 1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                 1675                 1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                 1690                 1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                 1705                 1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                 1720                 1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1730                 1735                 1740

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1745                 1750                 1755

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1760                 1765                 1770

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1775                 1780                 1785

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1790                 1795                 1800

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1805                 1810                 1815

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1820                 1825                 1830
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Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1835 1840 1845

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
1850 1855 1860

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
1865 1870 1875

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
1880 1885 1890

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1895 1900 1905

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1910 1915 1920

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1925 1930 1935

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
1940 1945 1950

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
1955 1960 1965

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
1970 1975 1980

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
1985 1990 1995

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2000 2005 2010

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2015 2020 2025

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Ala Thr Gly His His His His His His
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35 40 45

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
50 55 60

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
65 70 75 80

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
85 90 95

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
100 105 110

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
115 120 125

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
130 135 140

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
145 150 155 160

Ser Thr Glu Glu Gly Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro

210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
260 265 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
275 280 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
290 295 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305 310 315 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
325 330 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
340 345 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
355 360 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
370 375 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385 390 395 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
405 410 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
420 425 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
435 440 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
450 455 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465 470 475 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr

485 490 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
500 505 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
515 520 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
530 535 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545 550 555 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
565 570 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
580 585 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
595 600 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
610 615 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
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Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn

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Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
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Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
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Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
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Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
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Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
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Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
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Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
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Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
1070 1075 1080

Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln
1085 1090 1095

Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp
1100 1105 1110

Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala
1115 1120 1125

Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu
1130 1135 1140

Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
1145 1150 1155

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1160 1165 1170

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1175 1180 1185

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1190 1195 1200

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1205 1210 1215

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1220 1225 1230

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1235 1240 1245

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1250 1255 1260

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1265 1270 1275

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys

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Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
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Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
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Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
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Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
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Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1550 1555 1560

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1565 1570 1575

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1580 1585 1590

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1595 1600 1605

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1610 1615 1620

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1625 1630 1635

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Pro Gly Thr
1730 1735 1740

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1745 1750 1755

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
1760 1765 1770

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1775 1780 1785

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr

1790 1795 1800

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
1805 1810 1815

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
1820 1825 1830

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1835 1840 1845

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
1850 1855 1860

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
1865 1870 1875

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
1880 1885 1890

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1895 1900 1905

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1910 1915 1920

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1925 1930 1935

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
1940 1945 1950

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
1955 1960 1965

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
1970 1975 1980

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
1985 1990 1995

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2000 2005 2010

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Ala Thr Gly His His His His His His

2045 2050

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<211> 6162
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gctacctcag gcagtgagac gccaggaacg agcgagtccg ctacaccgga gagtgggcca 420

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<211> 2053
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Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
35 40 45

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
50 55 60

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
65 70 75 80

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
85 90 95

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
100 105 110

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
115 120 125

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
130 135 140

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
145 150 155 160

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ala Ser Ser
165 170 175

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Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
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Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445
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Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
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His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
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Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720
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Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990
```

```
Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245
```

```
Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500
```

```
Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1730                1735                1740

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1745                1750                1755
```

```
Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1760                 1765                 1770

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1775                 1780                 1785

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1790                 1795                 1800

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1805                 1810                 1815

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1820                 1825                 1830

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1835                 1840                 1845

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1850                 1855                 1860

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    1865                 1870                 1875

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    1880                 1885                 1890

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1895                 1900                 1905

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1910                 1915                 1920

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1925                 1930                 1935

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1940                 1945                 1950

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    1955                 1960                 1965

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    1970                 1975                 1980

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Glu Pro Ala
    1985                 1990                 1995

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2000                 2005                 2010
```

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2015 2020 2025

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
2030 2035 2040

Pro Ala Thr Gly His His His His His His
2045 2050

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<211> 6162
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<220>
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20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Gly Ala Pro Thr Ser Glu Ser Ala
35 40 45

Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu
50 55 60

Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
65 70 75 80

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala
85 90 95

Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
100 105 110

Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr
115 120 125

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Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr
    130                 135                 140

Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
145                 150                 155                 160

Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser
                165                 170                 175

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ala Ser Ser Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400
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Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670
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Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
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Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
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Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
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Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
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Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
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Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
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Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
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Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
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Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
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Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
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```

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Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
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Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
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Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                 1240                 1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                 1255                 1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                 1270                 1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                 1285                 1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                 1300                 1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                 1315                 1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                 1330                 1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                 1345                 1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                 1360                 1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                 1375                 1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                 1390                 1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                 1405                 1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                 1420                 1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                 1435                 1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                 1450                 1455
```

```
Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                 1465                 1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                 1480                 1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                 1495                 1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                 1510                 1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                 1525                 1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                 1540                 1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                 1555                 1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                 1570                 1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                 1585                 1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                 1600                 1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                 1615                 1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                 1630                 1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                 1645                 1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                 1660                 1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                 1675                 1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                 1690                 1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
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Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
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Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1730                1735                1740

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1745                1750                1755

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1760                1765                1770

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1775                1780                1785

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1790                1795                1800

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1805                1810                1815

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1820                1825                1830

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
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Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    1865                1870                1875

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    1880                1885                1890

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1895                1900                1905

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1910                1915                1920

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1925                1930                1935

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1940                1945                1950

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    1955                1960                1965
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Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
1970 1975 1980

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
1985 1990 1995

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2000 2005 2010

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2015 2020 2025

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Ala Thr Gly His His His His His His
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<211> 6162
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ggacccggaa ccagtaccga acctagcgag ggctctgctc cgggcagccc agccggctct 360

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20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

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Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350
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Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
                405                 410                 415

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser
            420                 425                 430

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly
        435                 440                 445

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
    450                 455                 460

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala
465                 470                 475                 480

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly
                485                 490                 495

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
            500                 505                 510

Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala
        515                 520                 525

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
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Ser Ser Thr Gly Ser Pro Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620
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Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895
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Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
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Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
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Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155
```

```
Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
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Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410
```

```
Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665
```

```
Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1730                1735                1740

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1745                1750                1755

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1760                1765                1770

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1775                1780                1785

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1790                1795                1800

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1805                1810                1815

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1820                1825                1830

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1835                1840                1845

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1850                1855                1860

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    1865                1870                1875

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    1880                1885                1890

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1895                1900                1905

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1910                1915                1920
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Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1925 1930 1935

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
1940 1945 1950

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
1955 1960 1965

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
1970 1975 1980

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
1985 1990 1995

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2000 2005 2010

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2015 2020 2025

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
2030 2035 2040

Pro Ala Thr Gly His His His His His His
2045 2050

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<211> 6162
<212> DNA
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<220>
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catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

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tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

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atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

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<211> 2053
<212> PRT
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20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys

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        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
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305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Gly
385 390 395 400

Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro
405 410 415

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
420 425 430

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
435 440 445

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
450 455 460

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
465 470 475 480

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
485 490 495

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
500 505 510

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
515 520 525

Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
530 535 540

Pro Gly Ala Ser Ser Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545 550 555 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
565 570 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile

580 585 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
595 600 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
610 615 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
645 650 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
835 840 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn

850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
885 890 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
900 905 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
915 920 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
930 935 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945 950 955 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
965 970 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
980 985 990

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
995 1000 1005

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser
1010 1015 1020

Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
1025 1030 1035

Ser Ala Pro Gly Ala Ser Ser Pro Pro Val Leu Lys Arg His Gln
1040 1045 1050

Ala Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
1055 1060 1065

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
1070 1075 1080

Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln
1085 1090 1095

Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp
1100 1105 1110

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1115 1120 1125

Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu
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Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
1145 1150 1155

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1160 1165 1170

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1175 1180 1185

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1190 1195 1200

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1205 1210 1215

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
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Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1235 1240 1245

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1250 1255 1260

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1265 1270 1275

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1280 1285 1290

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1295 1300 1305

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1310 1315 1320

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1325 1330 1335

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1340 1345 1350

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1355 1360 1365

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn

1370 1375 1380

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1385 1390 1395

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1400 1405 1410

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1415 1420 1425

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1430 1435 1440

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1445 1450 1455

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1460 1465 1470

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1475 1480 1485

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1490 1495 1500

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1505 1510 1515

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1520 1525 1530

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1535 1540 1545

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1550 1555 1560

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1565 1570 1575

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1580 1585 1590

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1595 1600 1605

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1610 1615 1620

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val

1625 1630 1635

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Pro Gly Thr
1730 1735 1740

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1745 1750 1755

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
1760 1765 1770

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1775 1780 1785

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1790 1795 1800

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
1805 1810 1815

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
1820 1825 1830

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1835 1840 1845

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
1850 1855 1860

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
1865 1870 1875

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu

1880 1885 1890

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1895 1900 1905

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1910 1915 1920

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1925 1930 1935

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
1940 1945 1950

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
1955 1960 1965

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
1970 1975 1980

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
1985 1990 1995

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2000 2005 2010

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2015 2020 2025

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Ala Thr Gly His His His His His His
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<211> 6162
<212> DNA
<213> Artificial Sequence

<220>
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atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

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<211> 2053
<212> PRT
<213> Artificial Sequence

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20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

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Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
                405                 410                 415

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser
            420                 425                 430

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly
        435                 440                 445

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
    450                 455                 460

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala
465                 470                 475                 480

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly
                485                 490                 495

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
            500                 505                 510

Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala
        515                 520                 525

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
    530                 535                 540
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Ser Ser Thr Gly Ser Pro Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815
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Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
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Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
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Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080
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Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                 1105                 1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                 1120                 1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                 1135                 1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                 1150                 1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                 1165                 1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                 1180                 1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                 1195                 1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                 1210                 1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                 1225                 1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                 1240                 1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                 1255                 1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                 1270                 1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                 1285                 1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                 1300                 1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                 1315                 1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                 1330                 1335
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```
Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
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Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590
```

```
Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1730                1735                1740

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1745                1750                1755

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1760                1765                1770

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1775                1780                1785

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1790                1795                1800

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1805                1810                1815

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1820                1825                1830

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1835                1840                1845
```

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
1850 1855 1860

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
1865 1870 1875

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
1880 1885 1890

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1895 1900 1905

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1910 1915 1920

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1925 1930 1935

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
1940 1945 1950

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
1955 1960 1965

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
1970 1975 1980

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
1985 1990 1995

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2000 2005 2010

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2015 2020 2025

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Ala Thr Gly His His His His His His
2045 2050

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<211> 6162
<212> DNA
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<220>
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<211> 2053
<212> PRT
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<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

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Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Gly
385                 390                 395                 400

Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro
                405                 410                 415

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
            420                 425                 430

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
        435                 440                 445

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
    450                 455                 460

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
465                 470                 475                 480

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
                485                 490                 495
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Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
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Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
        515                 520                 525

Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    530                 535                 540

Pro Gly Ala Ser Ser Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765
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Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
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Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035
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Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
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Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290
```

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Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
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Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                 1315                 1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                 1330                 1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                 1345                 1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                 1360                 1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                 1375                 1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                 1390                 1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                 1405                 1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                 1420                 1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                 1435                 1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                 1450                 1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                 1465                 1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                 1480                 1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                 1495                 1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                 1510                 1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                 1525                 1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                 1540                 1545
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Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1730                1735                1740

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1745                1750                1755

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1760                1765                1770

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1775                1780                1785

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1790                1795                1800
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Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1805                 1810                 1815

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1820                 1825                 1830

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1835                 1840                 1845

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1850                 1855                 1860

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    1865                 1870                 1875

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    1880                 1885                 1890

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1895                 1900                 1905

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1910                 1915                 1920

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1925                 1930                 1935

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1940                 1945                 1950

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    1955                 1960                 1965

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    1970                 1975                 1980

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Glu Pro Ala
    1985                 1990                 1995

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2000                 2005                 2010

Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    2015                 2020                 2025

Pro Ala  Ser Ser Glu Gln Lys  Leu Ile Ser Glu Glu  Asp Leu Ser
    2030                 2035                 2040

Pro Ala  Thr Gly His His His  His His His
    2045                 2050
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<210> 1098
<211> 6162
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

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ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

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gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

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gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcggc 1200

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cctgggacaa gcacagaacc ttcggaaggg agtgcccctg gaacatccga atcggcaacc 1440

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gatctgtcac ctgcaaccgg tcatcatcac catcaccatt ga 6162

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<211> 2053
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

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Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445
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Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720
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Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
            740                 745                 750

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
        755                 760                 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
    770                 775                 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785                 790                 795                 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                805                 810                 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            820                 825                 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
        835                 840                 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    850                 855                 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865                 870                 875                 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Gly Ala Pro Thr Ser Glu Ser Ala Thr Pro Glu
            980                 985                 990
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Ser Gly Pro Gly Ser Glu Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly
        995                 1000                1005


Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro
    1010                1015                1020


Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr
    1025                1030                1035


Pro Glu  Ser Gly Pro Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    1040                1045                1050


Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    1055                1060                1065


Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala
    1070                1075                1080


Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro
    1085                1090                1095


Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1100                1105                1110


Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    1115                1120                1125


Ala Ser  Ser Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140


Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155


Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170


Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185


Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200


Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215


Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230


Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245
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Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
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Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500
```

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Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
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Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1730                1735                1740

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1745                1750                1755
```

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Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1760                1765                1770

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1775                1780                1785

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1790                1795                1800

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1805                1810                1815

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1820                1825                1830

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1835                1840                1845

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1850                1855                1860

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    1865                1870                1875

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    1880                1885                1890

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1895                1900                1905

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1910                1915                1920

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1925                1930                1935

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1940                1945                1950

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    1955                1960                1965

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    1970                1975                1980

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Glu Pro Ala
    1985                1990                1995

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2000                2005                2010
```

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2015 2020 2025

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
2030 2035 2040

Pro Ala Thr Gly His His His His His His
2045 2050

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<211> 6162
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atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

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atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

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<211> 2056
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20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn

130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro

405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp

675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
740 745 750

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
755 760 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
770 775 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785 790 795 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
805 810 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
820 825 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
835 840 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
850 855 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865 870 875 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
885 890 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
900 905 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
915 920 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
930 935 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu

945 950 955 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
965 970 975

Arg Ala Gln Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
980 985 990

Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
995 1000 1005

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
1010 1015 1020

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
1025 1030 1035

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
1040 1045 1050

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
1055 1060 1065

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
1070 1075 1080

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
1085 1090 1095

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
1100 1105 1110

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser
1115 1120 1125

Ser Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val
1130 1135 1140

Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg
1145 1150 1155

Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
1160 1165 1170

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala
1175 1180 1185

Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu
1190 1195 1200

Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro

1205 1210 1215

Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala
1220 1225 1230

Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser
1235 1240 1245

Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
1250 1255 1260

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg
1265 1270 1275

Gln Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp
1280 1285 1290

Glu Thr Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys
1295 1300 1305

Arg Ala Pro Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu
1310 1315 1320

Asn Tyr Arg Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu
1325 1330 1335

Pro Gly Leu Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu
1340 1345 1350

Leu Ser Met Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser
1355 1360 1365

Gly His Val Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala
1370 1375 1380

Leu Tyr Asn Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu
1385 1390 1395

Pro Ser Lys Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu
1400 1405 1410

His Leu His Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn
1415 1420 1425

Lys Cys Gln Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp
1430 1435 1440

Phe Gln Ile Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys
1445 1450 1455

Leu Ala Arg Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr

1460 1465 1470

Lys Glu Pro Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met
1475 1480 1485

Ile Ile His Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser
1490 1495 1500

Ser Leu Tyr Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly
1505 1510 1515

Lys Lys Trp Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met
1520 1525 1530

Val Phe Phe Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile
1535 1540 1545

Phe Asn Pro Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr
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His Tyr Ser Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys
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Asp Leu Asn Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala
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Ile Ser Asp Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met
1595 1600 1605

Phe Ala Thr Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly
1610 1615 1620

Arg Ser Asn Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp
1625 1630 1635

Leu Gln Val Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr
1640 1645 1650

Thr Gln Gly Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu
1655 1660 1665

Phe Leu Ile Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe
1670 1675 1680

Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser
1685 1690 1695

Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg
1700 1705 1710

Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu

```
    1715                1720                1725

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    1730                1735                1740

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1745                1750                1755

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    1760                1765                1770

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1775                1780                1785

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1790                1795                1800

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    1805                1810                1815

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    1820                1825                1830

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1835                1840                1845

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1850                1855                1860

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser
    1865                1870                1875

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Thr Glu
    1880                1885                1890

Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1895                1900                1905

Glu Ser  Gly Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1910                1915                1920

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1925                1930                1935

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Ser  Glu Pro Ala
    1940                1945                1950

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Pro Ala Gly  Ser Pro Thr
    1955                1960                1965

Ser Thr  Glu Glu Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
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1970 1975 1980

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
1985 1990 1995

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
2000 2005 2010

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
2015 2020 2025

Gly Ser Ala Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu
2030 2035 2040

Asp Leu Ser Pro Ala Thr Gly His His His His His His
2045 2050 2055

<210> 1102
<211> 6171
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

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atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

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gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

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gccacaagtg gcagtgagac acccggcact tcagagagtg ccacccccga gagtggccca 6060

ggcactagta ccgagccctc tgaaggcagt gcgccagcct cgagcgaaca aaaactcatc 6120

tcagaagagg atctgtcacc tgcaaccggt catcatcacc atcaccattg a 6171

<210> 1103
<211> 2053
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1103
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

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Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365
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Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640
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Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
            740                 745                 750

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
        755                 760                 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
    770                 775                 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785                 790                 795                 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                805                 810                 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            820                 825                 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
        835                 840                 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    850                 855                 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865                 870                 875                 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910
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Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Gly Ala Pro Thr Ser Glu Ser Ala Thr
                965                 970                 975

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
            980                 985                 990

Pro Gly Thr Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
        995                 1000                1005

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1010                1015                1020

Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1040                1045                1050

Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr Glu Glu  Gly Ser Pro
    1055                1060                1065

Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Ser Pro  Ala Gly Ser
    1070                1075                1080

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1085                1090                1095

Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1100                1105                1110

Gly Ala  Ser Ser Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170
```

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Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425
```

```
Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680
```

```
Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1730                1735                1740

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1745                1750                1755

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1760                1765                1770

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1775                1780                1785

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1790                1795                1800

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1805                1810                1815

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1820                1825                1830

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1835                1840                1845

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1850                1855                1860

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    1865                1870                1875

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    1880                1885                1890

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1895                1900                1905

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1910                1915                1920

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1925                1930                1935
```

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
1940 1945 1950

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
1955 1960 1965

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
1970 1975 1980

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
1985 1990 1995

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2000 2005 2010

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2015 2020 2025

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
2030 2035 2040

Pro Ala Thr Gly His His His His His His
2045 2050

<210> 1104
<211> 6162
<212> DNA
<213> Artificial Sequence

<220>
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atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

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atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

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gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

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gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

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caaagaggaa accagataat gtcagacaag aggaatgtca tcctgttttc tgtatttgat 1740

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cccggtacct ccactgagcc cagcgagggg agtgcaccag gtgcctcgag cagtagctcc 3360

ccacatgttc taagaaacag ggctcagagt ggcagtgtcc ctcagttcaa gaaagttgtt 3420

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<211> 2056
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

```
Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320
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```
Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590
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Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
            740                 745                 750

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
        755                 760                 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
    770                 775                 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785                 790                 795                 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                805                 810                 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            820                 825                 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
        835                 840                 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    850                 855                 860
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Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865                 870                 875                 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
            980                 985                 990

Ser Pro Gly Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
        995                 1000                1005

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1010                1015                1020

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1025                1030                1035

Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1040                1045                1050

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1055                1060                1065

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1070                1075                1080

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1085                1090                1095

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1100                1105                1110

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
    1115                1120                1125
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Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1130                1135                1140

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1145                1150                1155

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1160                1165                1170

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1175                1180                1185

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1190                1195                1200

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1205                1210                1215

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1220                1225                1230

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1235                1240                1245

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1250                1255                1260

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1265                1270                1275

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1280                1285                1290

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1295                1300                1305

Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1310                1315                1320

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1325                1330                1335

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1340                1345                1350

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1355                1360                1365

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1370                1375                1380
```

```
Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1385                1390                1395

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1400                1405                1410

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1415                1420                1425

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1430                1435                1440

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1445                1450                1455

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1460                1465                1470

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1475                1480                1485

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1490                1495                1500

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1505                1510                1515

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1520                1525                1530

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1535                1540                1545

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1550                1555                1560

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1565                1570                1575

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1580                1585                1590

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1595                1600                1605

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1610                1615                1620

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1625                1630                1635
```

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Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1640                1645                1650

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1655                1660                1665

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1670                1675                1680

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1685                1690                1695

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    1700                1705                1710

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    1715                1720                1725

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    1730                1735                1740

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1745                1750                1755

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    1760                1765                1770

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1775                1780                1785

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1790                1795                1800

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    1805                1810                1815

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    1820                1825                1830

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1835                1840                1845

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1850                1855                1860

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser
    1865                1870                1875

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Thr Glu
    1880                1885                1890
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Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
1895 1900 1905

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1910 1915 1920

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
1925 1930 1935

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala
1940 1945 1950

Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
1955 1960 1965

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
1970 1975 1980

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
1985 1990 1995

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
2000 2005 2010

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
2015 2020 2025

Gly Ser Ala Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu
2030 2035 2040

Asp Leu Ser Pro Ala Thr Gly His His His His His His
2045 2050 2055

<210> 1106
<211> 6171
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

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<210> 1107
<211> 2053
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1107

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270
```

```
Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540
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Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
            740                 745                 750

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
        755                 760                 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
    770                 775                 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785                 790                 795                 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                805                 810                 815
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Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            820                 825                 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
        835                 840                 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    850                 855                 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865                 870                 875                 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Gly Ala Pro Thr Ser Glu Ser Ala Thr
                965                 970                 975

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
            980                 985                 990

Pro Gly Thr Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
        995                 1000                1005

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1010                1015                1020

Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1040                1045                1050

Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr Glu Glu  Gly Ser Pro
    1055                1060                1065

Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Ser Pro  Ala Gly Ser
    1070                1075                1080
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Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1085                1090                1095

Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1100                1105                1110

Gly Ala  Ser Ser Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335
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Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590
```

```
Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1730                1735                1740

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1745                1750                1755

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1760                1765                1770

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1775                1780                1785

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1790                1795                1800

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1805                1810                1815

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1820                1825                1830

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1835                1840                1845
```

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
1850 1855 1860

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
1865 1870 1875

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
1880 1885 1890

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1895 1900 1905

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1910 1915 1920

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1925 1930 1935

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
1940 1945 1950

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
1955 1960 1965

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
1970 1975 1980

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
1985 1990 1995

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2000 2005 2010

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2015 2020 2025

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
2030 2035 2040

Pro Ala Thr Gly His His His His His His
2045 2050

<210> 1108
<211> 6162
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

agagatctag cttcaggact cattggccct ctcctcatct gctacaaaga atctgtagat 1680

caaagaggaa accagataat gtcagacaag aggaatgtca tcctgttttc tgtatttgat 1740

gagaaccgaa gctggtacct cacagagaat atacaacgct ttctccccaa tccagctgga 1800

gtgcagcttg aggatccaga gttccaagcc tccaacatca tgcacagcat caatggctat 1860

gtttttgata gtttgcagtt gtcagtttgt ttgcatgagg tggcatactg gtacattcta 1920

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gacagttatg aagatatttc agcatacttg ctgagtaaaa acaatgccat tgaaccaaga 2220

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acctcagagt ccgccacccc cgagtccggc cctggctctg agcctgccac tagcggctcc 2340

gagactcctg gcacatccga aagcgctaca cccgagagtg gacccggcac ctctaccgag 2400

cccagtgagg gctccgcccc tggaacaagc accgagccca gcgaaggcag cgccccaggg 2460

acctccacag agcccagtga aggcagtgct cctggcacca gcaccgaacc aagcgagggc 2520

tctgcacccg ggacctccac cgagccaagc gaaggctctg cccctggcac ttccaccgag 2580

cccagcgaag gcagcgcccc tgggagcccc gctggctctc ccaccagcac tgaggagggc 2640

acatctaccg aaccaagtga aggctctgca ccaggtgcct cgagcccacc agtcttgaaa 2700

cgccatcaac gggaaataac tcgtactact cttcagtcag atcaagagga aatcgattat 2760

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aatcagagcc cccgcagctt tcaaaagaaa acacgacact attttattgc tgcagtggag 2880

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catttgggac tcctggggcc atatataaga gcagaagttg aagataatat catggtaact 3540

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gatcagaggc aaggagcaga acctagaaaa aactttgtca agcctaatga aaccaaaact 3660

tacttttgga aagtgcaaca tcatatggca cccactaaag atgagtttga ctgcaaagcc 3720

tgggcttatt tctctgatgt tgacctggaa aaagatgtgc actcaggcct gattggaccc 3780

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tatcgaggaa attccactgg aaccttaatg gtcttctttg gcaatgtgga ttcatctggg 4620

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aactctctag acccaccgtt actgactcgc taccttcgaa ttcaccccca gagttgggtg 5160

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ggcagtgaga cacccggcac ttcagagagt gccacccccg agagtggccc aggcactagt 6060

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gatctgtcac ctgcaaccgg tcatcatcac catcaccatt ga 6162

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<211> 2053
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg

225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys

500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
740 745 750

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
755 760 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly

770 775 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785 790 795 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
805 810 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
820 825 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
835 840 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
850 855 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865 870 875 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
885 890 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
900 905 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
915 920 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
930 935 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945 950 955 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
965 970 975

Arg Ala Gln Ser Gly Gly Ala Pro Thr Ser Glu Ser Ala Thr Pro Glu
980 985 990

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
995 1000 1005

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
1010 1015 1020

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
1025 1030 1035

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser

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    1040                1045                1050

Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    1055                1060                1065

Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala
    1070                1075                1080

Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro
    1085                1090                1095

Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1100                1105                1110

Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    1115                1120                1125

Ala Ser  Ser Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
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    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
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1550 1555 1560

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1565 1570 1575

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1580 1585 1590

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1595 1600 1605

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1610 1615 1620

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1625 1630 1635

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Pro Gly Thr
1730 1735 1740

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1745 1750 1755

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
1760 1765 1770

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1775 1780 1785

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1790 1795 1800

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly

1805 1810 1815

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
1820 1825 1830

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1835 1840 1845

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
1850 1855 1860

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
1865 1870 1875

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
1880 1885 1890

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1895 1900 1905

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1910 1915 1920

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1925 1930 1935

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
1940 1945 1950

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
1955 1960 1965

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
1970 1975 1980

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
1985 1990 1995

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2000 2005 2010

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2015 2020 2025

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
2030 2035 2040

Pro Ala Thr Gly His His His His His His
2045 2050

<210> 1110

<211> 6162
<212> DNA
<213> Artificial Sequence

<220>
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

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His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460
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Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735
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Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
            740                 745                 750

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
        755                 760                 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
    770                 775                 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785                 790                 795                 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                805                 810                 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            820                 825                 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
        835                 840                 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    850                 855                 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865                 870                 875                 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
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Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005
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Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
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Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1145                1150                1155

Pro Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    1160                1165                1170

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1175                1180                1185

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1190                1195                1200

Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1205                1210                1215

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1220                1225                1230

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Pro  Ala Gly Ser
    1235                1240                1245

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1250                1255                1260
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Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
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Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
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Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ala Ser  Ser Ala Pro
    1295                 1300                 1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                 1315                 1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                 1330                 1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                 1345                 1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                 1360                 1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                 1375                 1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                 1390                 1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                 1405                 1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                 1420                 1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                 1435                 1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                 1450                 1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                 1465                 1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                 1480                 1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                 1495                 1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                 1510                 1515
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Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
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Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
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Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
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Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
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Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
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Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1760                1765                1770
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Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
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Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
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Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
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Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
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Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
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Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
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Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
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Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
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Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    1955                 1960                 1965

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
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Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Glu Pro Ala
    1985                 1990                 1995

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2000                 2005                 2010

Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    2015                 2020                 2025
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Pro Ala Thr Gly His His His His His His
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20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

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Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415
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Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685
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Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
            740                 745                 750

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
        755                 760                 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
    770                 775                 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785                 790                 795                 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                805                 810                 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            820                 825                 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
        835                 840                 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    850                 855                 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865                 870                 875                 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960
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```
Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gly Ala Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    1160                1165                1170

Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1175                1180                1185

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1190                1195                1200

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala
    1205                1210                1215
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Ser Thr  Gly Thr Gly Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
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Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1235                 1240                 1245

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1250                 1255                 1260

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser
    1265                 1270                 1275

Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    1280                 1285                 1290

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1295                 1300                 1305

Ala Ser  Ser Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                 1315                 1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                 1330                 1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                 1345                 1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                 1360                 1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                 1375                 1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                 1390                 1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                 1405                 1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                 1420                 1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                 1435                 1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                 1450                 1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                 1465                 1470
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Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725
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Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
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Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1760                1765                1770

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1775                1780                1785

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
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Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1805                1810                1815

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
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Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
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Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    1865                1870                1875

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    1880                1885                1890

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1895                1900                1905

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1910                1915                1920

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1925                1930                1935

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1940                1945                1950

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    1955                1960                1965

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    1970                1975                1980
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Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
1985 1990 1995

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2000 2005 2010

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Ala Thr Gly His His His His His His
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

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Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365
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Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
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Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640
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Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
            740                 745                 750

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
        755                 760                 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
    770                 775                 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785                 790                 795                 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                805                 810                 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            820                 825                 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
        835                 840                 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    850                 855                 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865                 870                 875                 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910
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Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1145                1150                1155

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala
    1160                1165                1170
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Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
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Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
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Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
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Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala
    1220                1225                1230

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1235                1240                1245

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
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Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
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Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ala
    1280                1285                1290

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Ala Ser  Ser Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425
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Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
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Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
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Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
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Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
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Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
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Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
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Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1760                1765                1770

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
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Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
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Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
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Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
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Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
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Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1925                1930                1935
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Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
1955 1960 1965

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
1970 1975 1980

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
1985 1990 1995

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2000 2005 2010

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Ala Thr Gly His His His His His His
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<211> 2053
<212> PRT
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro

50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg

325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe

595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
740 745 750

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
755 760 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
770 775 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785 790 795 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
805 810 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
820 825 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
835 840 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
850 855 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly

865 870 875 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
885 890 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
900 905 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
915 920 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
930 935 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945 950 955 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
965 970 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
980 985 990

Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
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1010 1015 1020

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1025 1030 1035

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
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Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
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Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
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Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
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Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1100 1105 1110

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
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Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys

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Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Gly Ala
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1160 1165 1170

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1175 1180 1185

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Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Ala Pro
1295 1300 1305

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1310 1315 1320

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1325 1330 1335

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1340 1345 1350

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1355 1360 1365

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1370 1375 1380

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys

1385 1390 1395

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1400 1405 1410

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1415 1420 1425

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1430 1435 1440

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1445 1450 1455

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
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Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
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Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
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Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
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Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
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Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1535 1540 1545

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
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Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
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Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
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Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly

1640 1645 1650

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1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
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Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
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Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
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1760 1765 1770

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1775 1780 1785

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1790 1795 1800

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
1805 1810 1815

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
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Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1835 1840 1845

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
1850 1855 1860

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
1865 1870 1875

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
1880 1885 1890

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly

1895 1900 1905

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
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Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1925 1930 1935

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1985 1990 1995

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Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285
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Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560
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Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
            740                 745                 750

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
        755                 760                 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
    770                 775                 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785                 790                 795                 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                805                 810                 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            820                 825                 830
```

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Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
        835                 840                 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    850                 855                 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865                 870                 875                 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095
```

```
Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
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Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1145                1150                1155

Pro Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    1160                1165                1170

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1175                1180                1185

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1190                1195                1200

Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1205                1210                1215

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1220                1225                1230

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Pro  Ala Gly Ser
    1235                1240                1245

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1250                1255                1260

Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1265                1270                1275

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1280                1285                1290

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ala Ser  Ser Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350
```

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Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
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Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605
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Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1730                1735                1740

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1745                1750                1755

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1760                1765                1770

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1775                1780                1785

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1790                1795                1800

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1805                1810                1815

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1820                1825                1830

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1835                1840                1845

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1850                1855                1860
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Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
1865 1870 1875

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
1880 1885 1890

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1895 1900 1905

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1910 1915 1920

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1925 1930 1935

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
1940 1945 1950

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
1955 1960 1965

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
1970 1975 1980

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
1985 1990 1995

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2000 2005 2010

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Ala Thr Gly His His His His His His
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240
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Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510
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Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
            740                 745                 750

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
        755                 760                 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
    770                 775                 780
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```
Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785                 790                 795                 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                805                 810                 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            820                 825                 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
        835                 840                 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    850                 855                 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865                 870                 875                 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
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Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
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Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Gly Ala Pro Gly  Thr Pro Gly
    1160                1165                1170

Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1175                1180                1185

Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
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Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly
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Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly Pro Gly  Thr Pro Gly
    1220                1225                1230

Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1235                1240                1245

Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
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Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly
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Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr
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Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
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Ser Thr  Gly Ser Pro Ala Ser  Ser Thr Phe Lys Glu  Asn Tyr Arg
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Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
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Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
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Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
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Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
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Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
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Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
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Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
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Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
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Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
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Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
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Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1805                1810                1815
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Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
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Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
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Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
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Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
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Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1925                1930                1935

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1940                1945                1950

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    1955                1960                1965

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
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Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Glu Pro Ala
    1985                1990                1995

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2000                2005                2010

Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    2015                2020                2025

Pro Ala  Ser Ser Glu Gln Lys  Leu Ile Ser Glu Glu  Asp Leu Ser
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190
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His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
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Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
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Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
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Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
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Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735
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Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
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Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Gly Ala Pro Thr Ser Thr
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Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
    770                 775                 780

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
785                 790                 795                 800

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
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Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
            820                 825                 830

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
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Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr
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Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
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Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
                885                 890                 895

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser
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Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
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Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
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Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
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Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
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Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
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Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
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Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
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Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
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Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
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Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
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Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
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Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
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Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
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Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
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Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
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Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
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Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
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Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
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Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
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Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
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Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
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Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
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Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
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Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
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Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
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Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
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Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
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Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
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Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
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Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
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Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
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Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
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Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
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Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
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Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
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cagtgggccc caaagctggc cagacttcat tattccggat caatcaatgc ctggagcacc 3960

aaggagccct tttcttggat caaggtggat ctgttggcac caatgattat tcacggcatc 4020

aagacccagg gtgcccgtca gaagttctcc agcctctaca tctctcagtt tatcatcatg 4080

tatagtcttg atgggaagaa gtggcagact tatcgaggaa attccactgg aaccttaatg 4140

gtcttctttg gcaatgtgga ttcatctggg ataaaacaca atatttttaa ccctccaatt 4200

attgctcgat acatccgttt gcacccaact cattatagca ttcgcagcac tcttcgcatg 4260

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aatccaaaag agtggctgca agtggacttc cagaagacaa tgaaagtcac aggagtaact 4500

actcagggag taaaatctct gcttaccagc atgtatgtga aggagttcct catctccagc 4560

agtcaagatg gccatcagtg gactctcttt tttcagaatg gcaaagtaaa ggtttttcag 4620

ggaaatcaag actccttcac acctgtggtg aactctctag acccaccgtt actgactcgc 4680

taccttcgaa ttcaccccca gagttgggtg caccagattg ccctgaggat ggaggttctg 4740

ggctgcgagg cacaggacct ctacggcgcc ggatcacctg gggccgaaac ggccgaacaa 4800

aaactcatct cagaagagga tctgtcacct gcaaccggtc atcatcacca tcaccattga 4860

<210> 1125
<211> 1619
<212> PRT

<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 1125
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His

245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys

515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
740 745 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
755 760 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
770 775 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys

785 790 795 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
805 810 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
820 825 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
835 840 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
850 855 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865 870 875 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
885 890 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
900 905 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
915 920 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
930 935 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945 950 955 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
965 970 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
980 985 990

Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
995 1000 1005

Cys Asn Ile Gly Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr
1010 1015 1020

Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
1025 1030 1035

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
1040 1045 1050

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser

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    1055                1060                1065

Glu Gly  Ser Ala Pro Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    1070                1075                1080

Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    1085                1090                1095

Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr Pro Gly  Ser Glu Pro
    1100                1105                1110

Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Ser Pro Ala  Gly Ser Pro
    1115                1120                1125

Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1130                1135                1140

Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    1145                1150                1155

Ala Ser  Ser Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
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1310 1315 1320

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1325 1330 1335

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1340 1345 1350

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1355 1360 1365

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1370 1375 1380

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1385 1390 1395

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1400 1405 1410

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1415 1420 1425

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1430 1435 1440

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1445 1450 1455

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1460 1465 1470

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1475 1480 1485

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1490 1495 1500

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1505 1510 1515

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1520 1525 1530

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu

1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1580 1585 1590

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1595 1600 1605

Ser Pro Ala Thr Gly His His His His His His
1610 1615

<210> 1126
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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ctcggtgagc tgcctgtgga cgcaagattt cctcctagag tgccaaaatc ttttccattc 120

aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

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gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

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gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

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ggctgcgagg cacaggacct ctacggcgcc ggatcacctg gggccgaaac ggccgaacaa 4800

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<210> 1127
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300
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Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575
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Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845
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```
Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Gly Ala Pro Ser  Pro Ala Gly
    1010                1015                1020

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1025                1030                1035

Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    1040                1045                1050

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr
    1055                1060                1065

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Thr  Ser Thr Glu
    1070                1075                1080

Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1085                1090                1095

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1100                1105                1110
```

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Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Ser
    1115                1120                1125

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    1130                1135                1140

Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu
    1145                1150                1155

Gly Ser  Ala Pro Gly Ala Ser  Ser Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365
```

```
Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                1420                1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                1435                1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                1450                1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                1465                1470

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                1480                1485

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1490                1495                1500

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                1510                1515

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1520                1525                1530

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1535                1540                1545

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1550                1555                1560

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1565                1570                1575

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1580                1585                1590

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
    1595                1600                1605

Ser Pro  Ala Thr Gly His His  His His His His His
    1610                1615
```

<210> 1128
<211> 4860
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

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gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

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cttcagtcag atcaagagga aatcgattat gatgatacca tatcagttga aatgaagaag 2340

gaagattttg acatttatga tgaggatgaa aatcagagcc cccgcagctt tcaaaagaaa 2400

acacgacact attttattgc tgcagtggag aggctctggg attatgggat gagtagctcc 2460

ccacatgttc taagaaacag ggctcagagt ggcagtgtcc ctcagttcaa gaaagttgtt 2520

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ctcagcatgg gcagcaatga aaacatccat tctattcatt tcagtggaca tgtgttcact 3660

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ggctgcgagg cacaggacct ctacggcgcc ggatcacctg gggccgaaac ggccgaacaa 4800

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<210> 1129
<211> 1619
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1129

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

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Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350
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Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620
```

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Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895
```

```
Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
        995                 1000                1005

Pro Ser  Pro Ala Gly Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser
    1010                1015                1020

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro
    1025                1030                1035

Ser Glu  Gly Ser Ala Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1040                1045                1050

Thr Glu  Glu Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1055                1060                1065

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1070                1075                1080

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
    1085                1090                1095

Ser Gly  Ser Glu Thr Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1100                1105                1110

Glu Thr  Pro Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1115                1120                1125

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1130                1135                1140

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ala Ser  Ser Ala Pro
    1145                1150                1155
```

```
Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                1240                1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                1255                1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                1270                1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                1285                1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                1300                1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                1315                1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                1330                1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                1345                1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                1360                1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                1375                1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                1390                1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                1405                1410
```

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1415 1420 1425

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1430 1435 1440

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1445 1450 1455

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1460 1465 1470

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1475 1480 1485

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1490 1495 1500

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1505 1510 1515

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1520 1525 1530

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
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Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
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Ser Pro Ala Thr Gly His His His His His His
1610 1615

<210> 1130
<211> 4860
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<213> Artificial Sequence

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

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ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

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atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

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gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

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gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

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<223> Description of Artificial Sequence: Synthetic polypeptide

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

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Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
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Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400
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Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gly Ala Pro Thr Ser Thr Glu
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Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
            420                 425                 430

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
        435                 440                 445

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
    450                 455                 460

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
465                 470                 475                 480

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
                485                 490                 495

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
            500                 505                 510

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
        515                 520                 525

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
    530                 535                 540

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670
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Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940
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Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
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Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200
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Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
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Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                 1225                 1230

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                 1240                 1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                 1255                 1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                 1270                 1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                 1285                 1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                 1300                 1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                 1315                 1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                 1330                 1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                 1345                 1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                 1360                 1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                 1375                 1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                 1390                 1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                 1405                 1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                 1420                 1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                 1435                 1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                 1450                 1455
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Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
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Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                 1480                 1485

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1490                 1495                 1500

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                 1510                 1515

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1520                 1525                 1530

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1535                 1540                 1545

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1550                 1555                 1560

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1565                 1570                 1575

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
    1580                 1585                 1590

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
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Ser Pro  Ala Thr Gly His His  His His His His
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aatcagagcc cccgcagctt tcaaaagaaa acacgacact attttattgc tgcagtggag 2880

aggctctggg attatgggat gagtagctcc ccacatgttc taagaaacag ggctcagagt 2940

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<212> PRT
<213> Artificial Sequence

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu

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            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
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450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala

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Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
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Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
        755                 760                 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
    770                 775                 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785                 790                 795                 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                805                 810                 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            820                 825                 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
        835                 840                 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    850                 855                 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865                 870                 875                 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Gln
                885                 890                 895

Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala
            900                 905                 910

Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu
        915                 920                 925

Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val
    930                 935                 940

Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly
945                 950                 955                 960

Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu
                965                 970                 975

Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
            980                 985                 990

Tyr Ser Phe Tyr Ser Ser Leu Ile  Ser Tyr Glu Glu Asp  Gln Arg Gln
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Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr Lys
1010 1015 1020

Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp
1025 1030 1035

Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu
1040 1045 1050

Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val Cys
1055 1060 1065

His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr Val
1070 1075 1080

Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys Ser
1085 1090 1095

Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro Cys
1100 1105 1110

Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg Phe
1115 1120 1125

His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu Val
1130 1135 1140

Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met Gly
1145 1150 1155

Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val Phe
1160 1165 1170

Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn Leu
1175 1180 1185

Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys Ala
1190 1195 1200

Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His Ala
1205 1210 1215

Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln Thr
1220 1225 1230

Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile Thr
1235 1240 1245

Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg Leu

1250 1255 1260

His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro Phe
1265 1270 1275

Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His Gly
1280 1285 1290

Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr Ile
1295 1300 1305

Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp Gln
1310 1315 1320

Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe Gly
1325 1330 1335

Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro Pro
1340 1345 1350

Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser Ile
1355 1360 1365

Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn Ser
1370 1375 1380

Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp Ala
1385 1390 1395

Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp
1400 1405 1410

Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn Ala
1415 1420 1425

Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val Asp
1430 1435 1440

Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly Val
1445 1450 1455

Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile Ser
1460 1465 1470

Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn Gly
1475 1480 1485

Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro Val
1490 1495 1500

Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile

1505 1510 1515

His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val
1520 1525 1530

Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Pro Gly Thr Ser
1535 1540 1545

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr
1550 1555 1560

Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
1565 1570 1575

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
1580 1585 1590

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
1595 1600 1605

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser
1610 1615 1620

Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
1625 1630 1635

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
1640 1645 1650

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro
1655 1660 1665

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser
1670 1675 1680

Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
1685 1690 1695

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
1700 1705 1710

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
1715 1720 1725

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr
1730 1735 1740

Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
1745 1750 1755

Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu

1760 1765 1770

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
1775 1780 1785

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr
1790 1795 1800

Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
1805 1810 1815

Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
1820 1825 1830

Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro
1835 1840 1845

Ala Thr Gly His His His His His His
1850 1855

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<211> 5574
<212> DNA
<213> Artificial Sequence

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gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

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ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

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ccaggcacta gtaccgagcc ctctgaaggc agtgcgccag cctcgagcga acaaaaactc 5520

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<211> 1903
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

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His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430
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Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700
```

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Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
            740                 745                 750

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
        755                 760                 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
    770                 775                 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785                 790                 795                 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                805                 810                 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            820                 825                 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
        835                 840                 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    850                 855                 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865                 870                 875                 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Phe Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975
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```
Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1145                1150                1155

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1160                1165                1170

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1175                1180                1185

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1190                1195                1200

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1205                1210                1215

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1220                1225                1230
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```
Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1235                 1240                 1245

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1250                 1255                 1260

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1265                 1270                 1275

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1280                 1285                 1290

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1295                 1300                 1305

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1310                 1315                 1320

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1325                 1330                 1335

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1340                 1345                 1350

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1355                 1360                 1365

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1370                 1375                 1380

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1385                 1390                 1395

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1400                 1405                 1410

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1415                 1420                 1425

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1430                 1435                 1440

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1445                 1450                 1455

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1460                 1465                 1470

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1475                 1480                 1485
```

```
Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1490                1495                1500

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1505                1510                1515

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1520                1525                1530

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1535                1540                1545

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1550                1555                1560

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1565                1570                1575

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1580                1585                1590

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1595                1600                1605

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1610                1615                1620

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1625                1630                1635

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1640                1645                1650

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1655                1660                1665

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1670                1675                1680

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1685                1690                1695

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1700                1705                1710

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    1715                1720                1725

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    1730                1735                1740
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Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1745 1750 1755

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1760 1765 1770

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1775 1780 1785

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
1790 1795 1800

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
1805 1810 1815

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
1820 1825 1830

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
1835 1840 1845

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
1850 1855 1860

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
1865 1870 1875

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
1880 1885 1890

Pro Ala Thr Gly His His His His His His
1895 1900

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<211> 5712
<212> DNA
<213> Artificial Sequence

<220>
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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

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ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

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gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

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tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

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gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

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<211> 1903
<212> PRT
<213> Artificial Sequence

<220>
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20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro

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    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
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325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe

595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
740 745 750

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
755 760 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
770 775 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785 790 795 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
805 810 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
820 825 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
835 840 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
850 855 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly

865 870 875 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
885 890 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
900 905 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
915 920 925

Lys Lys Glu Asp Phe Asp Ile Phe Asp Glu Asp Glu Asn Gln Ser Pro
930 935 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945 950 955 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
965 970 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
980 985 990

Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
995 1000 1005

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1010 1015 1020

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1025 1030 1035

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1040 1045 1050

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1055 1060 1065

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1070 1075 1080

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1085 1090 1095

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1100 1105 1110

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1115 1120 1125

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys

1130 1135 1140

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1145 1150 1155

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1160 1165 1170

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1175 1180 1185

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1190 1195 1200

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1205 1210 1215

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1220 1225 1230

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1235 1240 1245

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1250 1255 1260

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1265 1270 1275

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1280 1285 1290

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1295 1300 1305

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1310 1315 1320

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1325 1330 1335

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1340 1345 1350

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1355 1360 1365

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1370 1375 1380

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro

1385 1390 1395

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1400 1405 1410

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1415 1420 1425

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1430 1435 1440

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1445 1450 1455

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1460 1465 1470

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1475 1480 1485

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1490 1495 1500

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1505 1510 1515

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1520 1525 1530

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1535 1540 1545

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1550 1555 1560

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1565 1570 1575

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Pro Gly Thr
1580 1585 1590

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1595 1600 1605

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
1610 1615 1620

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1625 1630 1635

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr

1640 1645 1650

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
1655 1660 1665

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
1670 1675 1680

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1685 1690 1695

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
1700 1705 1710

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
1715 1720 1725

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
1730 1735 1740

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1745 1750 1755

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1760 1765 1770

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1775 1780 1785

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
1790 1795 1800

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
1805 1810 1815

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
1820 1825 1830

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
1835 1840 1845

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
1850 1855 1860

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
1865 1870 1875

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Ala Thr Gly His His His His His His

1895 1900

<210> 1138
<211> 5712
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

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catcaccatt ga 5712

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<211> 1602
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

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Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495
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His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Thr Pro Gly
            740                 745                 750

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
        755                 760                 765
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Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    770                 775                 780

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
785                 790                 795                 800

Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala
                805                 810                 815

Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
            820                 825                 830

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
        835                 840                 845

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
    850                 855                 860

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
865                 870                 875                 880

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Ser
                885                 890                 895

Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys
            900                 905                 910

Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg
        915                 920                 925

Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg
    930                 935                 940

Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg
945                 950                 955                 960

Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu
                965                 970                 975

Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn
            980                 985                 990

Glu His Leu Gly Leu Leu Gly Pro  Tyr Ile Arg Ala Glu  Val Glu Asp
        995                 1000                1005

Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala Ser Arg  Pro Tyr Ser
    1010                1015                1020

Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu Asp Gln  Arg Gln Gly
    1025                1030                1035
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Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro Asn Glu  Thr Lys Thr
    1040                1045                1050

Tyr Phe  Trp Lys Val Gln His  His Met Ala Pro Thr  Lys Asp Glu
    1055                1060                1065

Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser Asp Val  Asp Leu Glu
    1070                1075                1080

Lys Asp  Val His Ser Gly Leu  Ile Gly Pro Leu Leu  Val Cys His
    1085                1090                1095

Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg Gln Val  Thr Val Gln
    1100                1105                1110

Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp Glu Thr  Lys Ser Trp
    1115                1120                1125

Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys Arg Ala  Pro Cys Asn
    1130                1135                1140

Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu Asn Tyr  Arg Phe His
    1145                1150                1155

Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu Pro Gly  Leu Val Met
    1160                1165                1170

Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu Leu Ser  Met Gly Ser
    1175                1180                1185

Asn Glu  Asn Ile His Ser Ile  His Phe Ser Gly His  Val Phe Thr
    1190                1195                1200

Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala Leu Tyr  Asn Leu Tyr
    1205                1210                1215

Pro Gly  Val Phe Glu Thr Val  Glu Met Leu Pro Ser  Lys Ala Gly
    1220                1225                1230

Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu His Leu  His Ala Gly
    1235                1240                1245

Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn Lys Cys  Gln Thr Pro
    1250                1255                1260

Leu Gly  Met Ala Ser Gly His  Ile Arg Asp Phe Gln  Ile Thr Ala
    1265                1270                1275

Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys Leu Ala  Arg Leu His
    1280                1285                1290
```

```
Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr Lys Glu  Pro Phe Ser
    1295                1300                1305

Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met Ile Ile  His Gly Ile
    1310                1315                1320

Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser Ser Leu  Tyr Ile Ser
    1325                1330                1335

Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly Lys Lys  Trp Gln Thr
    1340                1345                1350

Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met Val Phe  Phe Gly Asn
    1355                1360                1365

Val Asp  Ser Ser Gly Ile Lys  His Asn Ile Phe Asn  Pro Pro Ile
    1370                1375                1380

Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr His Tyr  Ser Ile Arg
    1385                1390                1395

Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys Asp Leu  Asn Ser Cys
    1400                1405                1410

Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala Ile Ser  Asp Ala Gln
    1415                1420                1425

Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met Phe Ala  Thr Trp Ser
    1430                1435                1440

Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly Arg Ser  Asn Ala Trp
    1445                1450                1455

Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp Leu Gln  Val Asp Phe
    1460                1465                1470

Gln Lys  Thr Met Lys Val Thr  Gly Val Thr Thr Gln  Gly Val Lys
    1475                1480                1485

Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu Phe Leu  Ile Ser Ser
    1490                1495                1500

Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe Phe Gln  Asn Gly Lys
    1505                1510                1515

Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser Phe Thr  Pro Val Val
    1520                1525                1530

Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg Tyr Leu  Arg Ile His
    1535                1540                1545
```

Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu
1550 1555 1560

Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala
1565 1570 1575

Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro
1580 1585 1590

Ala Thr Gly His His His His His His
1595 1600

<210> 1140
<211> 4809
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 1140
gccaccagaa gatactacct gggtgcagtg gaactgtcat gggactatat gcaaagtgat 60

ctcggtgagc tgcctgtgga cgcaagattt cctcctagag tgccaaaatc ttttccattc 120

aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

agagatctag cttcaggact cattggccct ctcctcatct gctacaaaga atctgtagat 1680

caaagaggaa accagataat gtcagacaag aggaatgtca tcctgttttc tgtatttgat 1740

gagaaccgaa gctggtacct cacagagaat atacaacgct ttctccccaa tccagctgga 1800

gtgcagcttg aggatccaga gttccaagcc tccaacatca tgcacagcat caatggctat 1860

gtttttgata gtttgcagtt gtcagtttgt ttgcatgagg tggcatactg gtacattcta 1920

agcattggag cacagactga cttcctttct gtcttcttct ctggatatac cttcaaacac 1980

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ggcatgaccg ccttactgaa ggtttctagt tgtgacaaga acactggtga ttattacgag 2160

gacagttatg aagatatttc agcatacttg ctgagtaaaa acaatgccat tgaaccaaga 2220

agcttctctc aaaacggcgc gccaggtaca cccggatcgg gtacagcgtc atcgagcccc 2280

ggtgcgtcac ctggtacgtc gagcacgggg tcgccagggg cgtcccctgg gacgtcctca 2340

acaggctcgc ccggtgcgtc acccggcacg tcgtccacgg gttcacctgg tagctcccct 2400

tccgcgtcca ctggcaccgg gcctggaact ccggggagcg gcacagcgag ctcgtcgccg 2460

ggagcatcgc ctgggacatc gagcaccggg tcgccaggag catcgcccgg aacatccagc 2520

acaggaagcc ccggcgcgtc gcccgggaca tcaagcacag gttccccggg atcgagcacg 2580

ccgtccggag ccactggatc accagggagc tcgacacctt ccggcgcaac gggatcgccc 2640

ggagccagcc cgggtacgtc aagcactggc tcccctgcct cgagctcaga tcaagaggaa 2700

atcgattatg atgataccat atcagttgaa atgaagaagg aagattttga catttatgat 2760

gaggatgaaa atcagagccc ccgcagcttt caaaagaaaa cacgacacta ttttattgct 2820

gcagtggaga ggctctggga ttatgggatg agtagctccc cacatgttct aagaaacagg 2880

gctcagagtg gcagtgtccc tcagttcaag aaagttgttt tccaggaatt tactgatggc 2940

tcctttactc agcccttata ccgtggagaa ctaaatgaac atttgggact cctggggcca 3000

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catatggcac ccactaaaga tgagtttgac tgcaaagcct gggcttattt ctctgatgtt 3240

gacctggaaa aagatgtgca ctcaggcctg attggacccc ttctggtctg ccacactaac 3300

acactgaacc ctgctcatgg gagacaagtg acagtacagg aatttgctct gtttttcacc 3360

atctttgatg agaccaaaag ctggtacttc actgaaaata tggaaagaaa ctgcagggct 3420

ccctgcaata tccagatgga agatcccact tttaaagaga attatcgctt ccatgcaatc 3480

aatggctaca taatggatac actacctggc ttagtaatgg ctcaggatca aaggattcga 3540

tggtatctgc tcagcatggg cagcaatgaa aacatccatt ctattcattt cagtggacat 3600

gtgttcactg tacgaaaaaa agaggagtat aaaatggcac tgtacaatct ctatccaggt 3660

gtttttgaga cagtggaaat gttaccatcc aaagctggaa tttggcgggt ggaatgcctt 3720

attggcgagc atctacatgc tgggatgagc acacttttc tggtgtacag caataagtgt 3780

cagactcccc tgggaatggc ttctggacac attagagatt ttcagattac agcttcagga 3840

caatatggac agtgggcccc aaagctggcc agacttcatt attccggatc aatcaatgcc 3900

tggagcacca aggagccctt ttcttggatc aaggtggatc tgttggcacc aatgattatt 3960

cacggcatca agacccaggg tgcccgtcag aagttctcca gcctctacat ctctcagttt 4020

atcatcatgt atagtcttga tgggaagaag tggcagactt atcgaggaaa ttccactgga 4080

accttaatgg tcttctttgg caatgtggat tcatctggga taaaacacaa tatttttaac 4140

cctccaatta ttgctcgata catccgtttg cacccaactc attatagcat tcgcagcact 4200

cttcgcatgg agttgatggg ctgtgattta aatagttgca gcatgccatt gggaatggag 4260

agtaaagcaa tatcagatgc acagattact gcttcatcct actttaccaa tatgtttgcc 4320

acctggtctc cttcaaaagc tcgacttcac ctccaaggga ggagtaatgc ctggagacct 4380

caggtgaata atccaaaaga gtggctgcaa gtggacttcc agaagacaat gaaagtcaca 4440

ggagtaacta ctcagggagt aaaatctctg cttaccagca tgtatgtgaa ggagttcctc 4500

atctccagca gtcaagatgg ccatcagtgg actctctttt ttcagaatgg caaagtaaag 4560

gtttttcagg gaaatcaaga ctccttcaca cctgtggtga actctctaga cccaccgtta 4620

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gaggttctgg gctgcgaggc acaggacctc tacggcgccg gatcacctgg ggccgaaacg 4740

gccgaacaaa aactcatctc agaagaggat ctgtcacctg caaccggtca tcatcaccat 4800

caccattga 4809

<210> 1141
<211> 1886
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1141
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro

```
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
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290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe

565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Thr Pro Gly
740 745 750

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
755 760 765

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
770 775 780

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
785 790 795 800

Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala
805 810 815

Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
820 825 830

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro

835 840 845

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
850 855 860

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
865 870 875 880

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Ser
885 890 895

Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys
900 905 910

Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg
915 920 925

Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg
930 935 940

Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg
945 950 955 960

Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu
965 970 975

Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn
980 985 990

Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp
995 1000 1005

Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser
1010 1015 1020

Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg Gln Gly
1025 1030 1035

Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr Lys Thr
1040 1045 1050

Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp Glu
1055 1060 1065

Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu
1070 1075 1080

Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val Cys His
1085 1090 1095

Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr Val Gln

1100 1105 1110

Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys Ser Trp
1115 1120 1125

Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro Cys Asn
1130 1135 1140

Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg Phe His
1145 1150 1155

Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu Val Met
1160 1165 1170

Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met Gly Ser
1175 1180 1185

Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val Phe Thr
1190 1195 1200

Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn Leu Tyr
1205 1210 1215

Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys Ala Gly
1220 1225 1230

Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His Ala Gly
1235 1240 1245

Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln Thr Pro
1250 1255 1260

Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile Thr Ala
1265 1270 1275

Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg Leu His
1280 1285 1290

Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro Phe Ser
1295 1300 1305

Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His Gly Ile
1310 1315 1320

Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr Ile Ser
1325 1330 1335

Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp Gln Thr
1340 1345 1350

Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe Gly Asn

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    1355                1360                1365

Val Asp  Ser Ser Gly Ile Lys  His Asn Ile Phe Asn  Pro Pro Ile
    1370                1375                1380

Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr His Tyr  Ser Ile Arg
    1385                1390                1395

Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys Asp Leu  Asn Ser Cys
    1400                1405                1410

Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala Ile Ser  Asp Ala Gln
    1415                1420                1425

Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met Phe Ala  Thr Trp Ser
    1430                1435                1440

Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly Arg Ser  Asn Ala Trp
    1445                1450                1455

Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp Leu Gln  Val Asp Phe
    1460                1465                1470

Gln Lys  Thr Met Lys Val Thr  Gly Val Thr Thr Gln  Gly Val Lys
    1475                1480                1485

Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu Phe Leu  Ile Ser Ser
    1490                1495                1500

Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe Phe Gln  Asn Gly Lys
    1505                1510                1515

Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser Phe Thr  Pro Val Val
    1520                1525                1530

Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg Tyr Leu  Arg Ile His
    1535                1540                1545

Pro Gln  Ser Trp Val His Gln  Ile Ala Leu Arg Met  Glu Val Leu
    1550                1555                1560

Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala Pro Gly  Thr Ser Glu
    1565                1570                1575

Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser
    1580                1585                1590

Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1595                1600                1605

Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly
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1610 1615 1620

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
1625 1630 1635

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro
1640 1645 1650

Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
1655 1660 1665

Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
1670 1675 1680

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
1685 1690 1695

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro
1700 1705 1710

Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
1715 1720 1725

Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
1730 1735 1740

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu
1745 1750 1755

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
1760 1765 1770

Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu
1775 1780 1785

Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
1790 1795 1800

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
1805 1810 1815

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser
1820 1825 1830

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
1835 1840 1845

Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Ala
1850 1855 1860

Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala

1865 1870 1875

Thr Gly His His His His His His
1880 1885

<210> 1142
<211> 5661
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 1142
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ctcggtgagc tgcctgtgga cgcaagattt cctcctagag tgccaaaatc ttttccattc 120

aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

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Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
                405                 410                 415

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            420                 425                 430

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
        435                 440                 445

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    450                 455                 460

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
465                 470                 475                 480

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
                485                 490                 495
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Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
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Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
    530                 535                 540

Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765
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Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
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Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
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Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
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Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
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Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
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Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290
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Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
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Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
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Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                 1330                 1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                 1345                 1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                 1360                 1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                 1375                 1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                 1390                 1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                 1405                 1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
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Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                 1435                 1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                 1450                 1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                 1465                 1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                 1480                 1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                 1495                 1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                 1510                 1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                 1525                 1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                 1540                 1545
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Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
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Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
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Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
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Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1760                1765                1770

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
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Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
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Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
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Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
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Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
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Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    1880                 1885                 1890

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1895                 1900                 1905

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1925                 1930                 1935

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1940                 1945                 1950

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    1955                 1960                 1965

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    1970                 1975                 1980

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Glu Pro Ala
    1985                 1990                 1995

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2000                 2005                 2010

Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    2015                 2020                 2025

Pro Ala  Ser Ser Glu Gln Lys  Leu Ile Ser Glu Glu  Asp Leu Ser
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Pro Ala  Thr Gly His His His  His His His
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<211> 2053
<212> PRT
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<220>
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<400> 1145

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

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Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gly Ala Pro Thr Ser Thr Glu
                405                 410                 415

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
            420                 425                 430

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
        435                 440                 445
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Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
    450                 455                 460

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
465                 470                 475                 480

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
                485                 490                 495

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
            500                 505                 510

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
        515                 520                 525

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
    530                 535                 540

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720
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Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990
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Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245
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Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500
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Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1745                1750                1755
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Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1760                 1765                 1770

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1775                 1780                 1785

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1790                 1795                 1800

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1805                 1810                 1815

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1820                 1825                 1830

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1835                 1840                 1845

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1850                 1855                 1860

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    1865                 1870                 1875

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    1880                 1885                 1890

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1895                 1900                 1905

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1910                 1915                 1920

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1925                 1930                 1935

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1940                 1945                 1950

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    1955                 1960                 1965

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    1970                 1975                 1980

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Glu Pro Ala
    1985                 1990                 1995

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2000                 2005                 2010
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2015 2020 2025

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Ala Thr Gly His His His His His His
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

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Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
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Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
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Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400
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Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
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Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
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Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
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Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
465                 470                 475                 480

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
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Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
           500                 505                 510

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
       515                 520                 525

Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
   530                 535                 540

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
               565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
           580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
       595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
   610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
               645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
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Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
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Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
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Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
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Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
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Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
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Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
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Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
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Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
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Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
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Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
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Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
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Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
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Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
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Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
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Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1010 1015 1020

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
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Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
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Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
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Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
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Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
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Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
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Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
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Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
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Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
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Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
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Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
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Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
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Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
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Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
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Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
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Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
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Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
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Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
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Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
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Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
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Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
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Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
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Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
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Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
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Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
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Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
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Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
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Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
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Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
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Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Gly Ser Pro
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35 40 45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
50 55 60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65 70 75 80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
85 90 95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
100 105 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
115 120 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
130 135 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145 150 155 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr

180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
260 265 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
275 280 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
290 295 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
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Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
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Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
355 360 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
370 375 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385 390 395 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
405 410 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
420 425 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
435 440 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser

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His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
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Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
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Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu
545                 550                 555                 560

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
                565                 570                 575

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            580                 585                 590

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
        595                 600                 605

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    610                 615                 620

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
625                 630                 635                 640

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
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Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
            660                 665                 670

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
        675                 680                 685

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp
    690                 695                 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
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725 730 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
740 745 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
755 760 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
770 775 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785 790 795 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
805 810 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
820 825 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
835 840 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
850 855 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865 870 875 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
885 890 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
900 905 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
915 920 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
930 935 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945 950 955 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
965 970 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
980 985 990

His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val

995 1000 1005

Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr
1010 1015 1020

Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu
1025 1030 1035

Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His Gln
1040 1045 1050

Arg Glu Ile Thr Arg Thr Thr Leu Gln Gly Ala Pro Gly Thr Pro
1055 1060 1065

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr
1070 1075 1080

Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
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1175 1180 1185

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Ser Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Gln Glu Glu Ile
1205 1210 1215

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
1220 1225 1230

Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln
1235 1240 1245

Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp

1250 1255 1260

Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala
1265 1270 1275

Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu
1280 1285 1290

Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
1295 1300 1305

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
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Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1325 1330 1335

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
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Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
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Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1385 1390 1395

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1400 1405 1410

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1415 1420 1425

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1430 1435 1440

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1445 1450 1455

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1460 1465 1470

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1475 1480 1485

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1490 1495 1500

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val

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Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1520 1525 1530

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
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Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1550 1555 1560

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1565 1570 1575

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1580 1585 1590

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1595 1600 1605

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1610 1615 1620

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
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Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1640 1645 1650

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1655 1660 1665

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1670 1675 1680

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1685 1690 1695

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
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Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1715 1720 1725

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1730 1735 1740

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1745 1750 1755

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn

1760 1765 1770

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1775 1780 1785

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1790 1795 1800

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1805 1810 1815

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1820 1825 1830

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1835 1840 1845

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1850 1855 1860

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1985 1990 1995

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Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly

2015 2020 2025

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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
35 40 45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
50 55 60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65 70 75 80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
85 90 95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
100 105 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
115 120 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
130 135 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145 150 155 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

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Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
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Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460
```

```
His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu
545                 550                 555                 560

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
                565                 570                 575

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            580                 585                 590

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
        595                 600                 605

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    610                 615                 620

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
625                 630                 635                 640

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
                645                 650                 655

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
            660                 665                 670

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
        675                 680                 685

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp
    690                 695                 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735
```

```
Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005
```

```
Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
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Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Gly  Ala Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1130                1135                1140

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1145                1150                1155

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1160                1165                1170

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1175                1180                1185

Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1190                1195                1200

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1205                1210                1215

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1220                1225                1230

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1235                1240                1245

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1250                1255                1260
```

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Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
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Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1280                 1285                 1290

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1295                 1300                 1305

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1310                 1315                 1320

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1325                 1330                 1335

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1340                 1345                 1350

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1355                 1360                 1365

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1370                 1375                 1380

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1385                 1390                 1395

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1400                 1405                 1410

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1415                 1420                 1425

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1430                 1435                 1440

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1445                 1450                 1455

Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1460                 1465                 1470

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1475                 1480                 1485

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1490                 1495                 1500

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1505                 1510                 1515
```

```
Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
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Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1535                1540                1545

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1550                1555                1560

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1565                1570                1575

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1580                1585                1590

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1595                1600                1605

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1610                1615                1620

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1625                1630                1635

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1640                1645                1650

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1655                1660                1665

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1670                1675                1680

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1685                1690                1695

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1700                1705                1710

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1715                1720                1725

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1730                1735                1740

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1745                1750                1755

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1760                1765                1770
```

```
Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
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Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1790                 1795                 1800

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1805                 1810                 1815

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1820                 1825                 1830

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1835                 1840                 1845

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    1850                 1855                 1860

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    1865                 1870                 1875

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    1880                 1885                 1890

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1895                 1900                 1905

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    1910                 1915                 1920

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1925                 1930                 1935

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1940                 1945                 1950

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    1955                 1960                 1965

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    1970                 1975                 1980

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1985                 1990                 1995

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2000                 2005                 2010

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser
    2015                 2020                 2025
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Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu
2030 2035 2040

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
2045 2050 2055

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2060 2065 2070

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2075 2080 2085

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala
2090 2095 2100

Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
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Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
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Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
2150 2155 2160

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
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Gly Ser Ala Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu
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Asp Leu Ser Pro Ala Thr Gly His His His His His His
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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
35 40 45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
50 55 60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65 70 75 80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
85 90 95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
100 105 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
115 120 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
130 135 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145 150 155 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

```
Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
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Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460
```

```
His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu
545                 550                 555                 560

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
                565                 570                 575

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            580                 585                 590

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
        595                 600                 605

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    610                 615                 620

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
625                 630                 635                 640

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
                645                 650                 655

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
            660                 665                 670

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
        675                 680                 685

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp
    690                 695                 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735
```

```
Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005
```

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Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
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Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                 1030                 1035

Pro Arg  Ser Phe Ser Gln Asn  Pro Pro Val Leu Lys  Arg His Gln
    1040                 1045                 1050

Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                 1060                 1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                 1075                 1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                 1090                 1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                 1105                 1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                 1120                 1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                 1135                 1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                 1150                 1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                 1165                 1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                 1180                 1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                 1195                 1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                 1210                 1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                 1225                 1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                 1240                 1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                 1255                 1260
```

```
Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1295                1300                1305

Pro Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    1310                1315                1320

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1325                1330                1335

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1340                1345                1350

Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1355                1360                1365

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1370                1375                1380

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Pro  Ala Gly Ser
    1385                1390                1395

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1400                1405                1410

Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1415                1420                1425

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1430                1435                1440

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ala Ser  Ser Ala Pro
    1445                1450                1455

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515
```

```
Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                1555                1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                1615                1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770
```

```
Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                1960                1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                1975                1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                1990                1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                2005                2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                2020                2025
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Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
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Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
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Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
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Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
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2090 2095 2100

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
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Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
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Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
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Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
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Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Ala Thr Gly His His His His His His
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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
35 40 45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
50 55 60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65 70 75 80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
85 90 95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
100 105 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
115 120 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
130 135 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145 150 155 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

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Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
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Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460
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His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735
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Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
        915                 920                 925

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
    930                 935                 940

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
945                 950                 955                 960

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
                965                 970                 975

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly
            980                 985                 990

Thr Ser Ser Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
        995                 1000                1005
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Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
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Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser
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Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    1040                 1045                 1050

Ser Ser  Thr Gly Ser Pro Ala  Ser Ser Ser Asp Gln  Glu Glu Ile
    1055                 1060                 1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                 1075                 1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                 1090                 1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                 1105                 1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                 1120                 1125

Gln Gly  Ala Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1130                 1135                 1140

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1145                 1150                 1155

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1160                 1165                 1170

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1175                 1180                 1185

Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1190                 1195                 1200

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1205                 1210                 1215

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1220                 1225                 1230

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1235                 1240                 1245

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1250                 1255                 1260
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Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
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Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1280                 1285                 1290

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1295                 1300                 1305

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1310                 1315                 1320

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1325                 1330                 1335

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1340                 1345                 1350

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1355                 1360                 1365

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1370                 1375                 1380

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1385                 1390                 1395

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1400                 1405                 1410

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1415                 1420                 1425

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1430                 1435                 1440

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1445                 1450                 1455

Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1460                 1465                 1470

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1475                 1480                 1485

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1490                 1495                 1500

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1505                 1510                 1515
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Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
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Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1535                1540                1545

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1550                1555                1560

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1565                1570                1575

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1580                1585                1590

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1595                1600                1605

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1610                1615                1620

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1625                1630                1635

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1640                1645                1650

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1655                1660                1665

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1670                1675                1680

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1685                1690                1695

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1700                1705                1710

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1715                1720                1725

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1730                1735                1740

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1745                1750                1755

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1760                1765                1770
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Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
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Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1790                 1795                 1800

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1805                 1810                 1815

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1820                 1825                 1830

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1835                 1840                 1845

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    1850                 1855                 1860

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    1865                 1870                 1875

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    1880                 1885                 1890

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1895                 1900                 1905

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    1910                 1915                 1920

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1925                 1930                 1935

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1940                 1945                 1950

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    1955                 1960                 1965

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    1970                 1975                 1980

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1985                 1990                 1995

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2000                 2005                 2010

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser
    2015                 2020                 2025
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Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu
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Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
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Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
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Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
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Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala
2090 2095 2100

Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
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Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
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Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
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Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
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Gly Ser Ala Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu
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35 40 45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
50 55 60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65 70 75 80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
85 90 95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
100 105 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
115 120 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
130 135 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145 150 155 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu

195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
260 265 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
275 280 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
290 295 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305 310 315 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
325 330 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
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Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
355 360 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
370 375 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385 390 395 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
405 410 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
420 425 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
435 440 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
450 455 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro

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Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
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Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
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740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
835 840 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
885 890 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
900 905 910

Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
915 920 925

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
930 935 940

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
945 950 955 960

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
965 970 975

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly
980 985 990

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
995 1000 1005

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro

1010 1015 1020

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
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Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
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Ser Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Gln Glu Glu Ile
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Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
1070 1075 1080

Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln
1085 1090 1095

Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp
1100 1105 1110

Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala
1115 1120 1125

Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu
1130 1135 1140

Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
1145 1150 1155

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1160 1165 1170

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1175 1180 1185

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1190 1195 1200

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
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Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1220 1225 1230

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1235 1240 1245

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1250 1255 1260

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr

1265 1270 1275

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1280 1285 1290

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Gly Ala
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1310 1315 1320

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala
1325 1330 1335

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1340 1345 1350

Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
1355 1360 1365

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
1370 1375 1380

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser
1385 1390 1395

Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser
1400 1405 1410

Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
1415 1420 1425

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
1430 1435 1440

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Ala Pro
1445 1450 1455

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1460 1465 1470

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1475 1480 1485

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1490 1495 1500

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
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Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn

1520 1525 1530

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1535 1540 1545

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1550 1555 1560

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1565 1570 1575

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1580 1585 1590

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1595 1600 1605

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
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Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
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Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1640 1645 1650

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1655 1660 1665

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1670 1675 1680

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1685 1690 1695

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
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Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1715 1720 1725

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1730 1735 1740

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1745 1750 1755

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1760 1765 1770

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val

1775 1780 1785

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
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Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1805 1810 1815

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1820 1825 1830

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1835 1840 1845

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1850 1855 1860

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1865 1870 1875

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Pro Gly Thr
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Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1940 1945 1950

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
1955 1960 1965

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Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1985 1990 1995

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2000 2005 2010

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
35 40 45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
50 55 60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65 70 75 80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
85 90 95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
100 105 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
115 120 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
130 135 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145 150 155 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

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Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480
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Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750
```

```
Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
            980                 985                 990

Ser Pro Gly Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
        995                 1000                1005

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
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Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
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Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1040                 1045                 1050

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1055                 1060                 1065

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1070                 1075                 1080

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1085                 1090                 1095

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1100                 1105                 1110

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
    1115                 1120                 1125

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1130                 1135                 1140

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1145                 1150                 1155

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1160                 1165                 1170

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1175                 1180                 1185

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1190                 1195                 1200

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1205                 1210                 1215

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1220                 1225                 1230

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1235                 1240                 1245

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1250                 1255                 1260

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1265                 1270                 1275
```

```
Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
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Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1295                1300                1305

Arg Gly  Ala Pro Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1310                1315                1320

Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser
    1325                1330                1335

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
    1340                1345                1350

Ser Gly  Ser Glu Thr Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
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Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
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Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Pro
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Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Ser Pro  Ala Gly Ser
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Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1415                1420                1425

Thr Glu  Glu Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1430                1435                1440

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Ala Ser
    1445                1450                1455

Ser Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1460                1465                1470

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1475                1480                1485

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1490                1495                1500

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1505                1510                1515

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1520                1525                1530
```

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Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1535                1540                1545

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1550                1555                1560

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1565                1570                1575

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1580                1585                1590

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1595                1600                1605

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1610                1615                1620

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1625                1630                1635

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1640                1645                1650

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1655                1660                1665

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1670                1675                1680

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1685                1690                1695

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1700                1705                1710

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1715                1720                1725

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1730                1735                1740

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1745                1750                1755

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1760                1765                1770

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1775                1780                1785
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Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1790                 1795                 1800

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1805                 1810                 1815

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1820                 1825                 1830

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1835                 1840                 1845

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    1850                 1855                 1860

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    1865                 1870                 1875

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    1880                 1885                 1890

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1895                 1900                 1905

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
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Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
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Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1940                 1945                 1950

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    1955                 1960                 1965

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
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Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1985                 1990                 1995

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2000                 2005                 2010

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser
    2015                 2020                 2025

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Thr Glu
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Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
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Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
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Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
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Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
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Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
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Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
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Gly Ser Ala Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu
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Asp Leu Ser Pro Ala Thr Gly His His His His His His
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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
35 40 45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
50 55 60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65 70 75 80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
85 90 95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
100 105 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
115 120 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
130 135 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145 150 155 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

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Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480
```

```
Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu
545                 550                 555                 560

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
                565                 570                 575

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            580                 585                 590

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
        595                 600                 605

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    610                 615                 620

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
625                 630                 635                 640

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
                645                 650                 655

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
            660                 665                 670

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
        675                 680                 685

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp
    690                 695                 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750
```

```
Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020
```

```
Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Gly Ala Pro  Gly Thr Pro
    1055                1060                1065

Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly Ala Ser  Pro Gly Thr
    1070                1075                1080

Ser Ser  Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
    1085                1090                1095

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1100                1105                1110

Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr Gly Pro  Gly Thr Pro
    1115                1120                1125

Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly Ala Ser  Pro Gly Thr
    1130                1135                1140

Ser Ser  Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
    1145                1150                1155

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1160                1165                1170

Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser
    1175                1180                1185

Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    1190                1195                1200

Ser Ser  Thr Gly Ser Pro Ala  Ser Ser Ser Asp Gln  Glu Glu Ile
    1205                1210                1215

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1220                1225                1230

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1235                1240                1245

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1250                1255                1260

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1265                1270                1275
```

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Gln Gly  Ala Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1280                1285                1290

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1295                1300                1305

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1310                1315                1320

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1325                1330                1335

Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1340                1345                1350

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1355                1360                1365

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1370                1375                1380

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1385                1390                1395

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1400                1405                1410

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
    1415                1420                1425

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1430                1435                1440

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1445                1450                1455

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1460                1465                1470

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1475                1480                1485

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1490                1495                1500

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1505                1510                1515

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1520                1525                1530
```

```
Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1535                1540                1545

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1550                1555                1560

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1565                1570                1575

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1580                1585                1590

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1595                1600                1605

Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1610                1615                1620

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1625                1630                1635

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1640                1645                1650

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1655                1660                1665

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1670                1675                1680

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1685                1690                1695

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1700                1705                1710

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1715                1720                1725

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1730                1735                1740

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1745                1750                1755

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1760                1765                1770

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1775                1780                1785
```

```
Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1790                 1795                 1800

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1805                 1810                 1815

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1820                 1825                 1830

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1835                 1840                 1845

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1850                 1855                 1860

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1865                 1870                 1875

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1880                 1885                 1890

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1895                 1900                 1905

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1910                 1915                 1920

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1925                 1930                 1935

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1940                 1945                 1950

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1955                 1960                 1965

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1970                 1975                 1980

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1985                 1990                 1995

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    2000                 2005                 2010

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    2015                 2020                 2025

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    2030                 2035                 2040
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2045                 2050                 2055

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    2060                 2065                 2070

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    2075                 2080                 2085

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2090                 2095                 2100

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    2105                 2110                 2115

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    2120                 2125                 2130

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    2135                 2140                 2145

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2150                 2155                 2160

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser
    2165                 2170                 2175

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Thr Glu
    2180                 2185                 2190

Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2195                 2200                 2205

Glu Ser  Gly Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2210                 2215                 2220

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2225                 2230                 2235

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Ser  Glu Pro Ala
    2240                 2245                 2250

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Pro Ala Gly  Ser Pro Thr
    2255                 2260                 2265

Ser Thr  Glu Glu Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    2270                 2275                 2280

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    2285                 2290                 2295
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Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
2300 2305 2310

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
2315 2320 2325

Gly Ser Ala Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu
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Asp Leu Ser Pro Ala Thr Gly His His His His His His
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acggggacgg gaccgggggc gtcaccagga acgtcctcca ccggctcgcc gggtgcatca 180

cccggaacgt catcgaccgg atcgccaggg agctcgacgc catcaggcgc aacaggatca 240

cctggctcaa gccctagcgc gtcaaccggc acgggtccgg gtgcctcccc tggcacgtcc 300

agcaccggat cacccggatc gagcccatcc gcctcaaccg gaaccggacc cggtacacca 360

gggtcgggaa cagcctcctc gtcaccaggc tcctcaaccc cctcgggagc cacgggttcg 420

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Met Gln Gly Ala Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
35 40 45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
50 55 60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65 70 75 80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
85 90 95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
100 105 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
115 120 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
130 135 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145 150 155 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
260 265 270

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Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540
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Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu
545                 550                 555                 560

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
                565                 570                 575

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            580                 585                 590

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
        595                 600                 605

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    610                 615                 620

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
625                 630                 635                 640

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
                645                 650                 655

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
            660                 665                 670

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
        675                 680                 685

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp
    690                 695                 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815
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Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020

Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Gly Ala Pro  Gly Thr Pro
    1055                1060                1065

Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly Ala Ser  Pro Gly Thr
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Ser Ser  Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
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Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1100                1105                1110

Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr Gly Pro  Gly Thr Pro
    1115                1120                1125

Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly Ala Ser  Pro Gly Thr
    1130                1135                1140

Ser Ser  Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
    1145                1150                1155

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1160                1165                1170

Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser
    1175                1180                1185

Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    1190                1195                1200

Ser Ser  Thr Gly Ser Pro Ala  Ser Ser Ser Asp Gln  Glu Glu Ile
    1205                1210                1215

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1220                1225                1230

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1235                1240                1245

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1250                1255                1260

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1265                1270                1275

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1280                1285                1290

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1295                1300                1305

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1310                1315                1320

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1325                1330                1335
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Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
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Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1355                1360                1365

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1370                1375                1380

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1385                1390                1395

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1400                1405                1410

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1415                1420                1425

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1430                1435                1440

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1445                1450                1455

Pro Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    1460                1465                1470

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1475                1480                1485

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1490                1495                1500

Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1505                1510                1515

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
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Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Pro  Ala Gly Ser
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Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
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Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
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Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1580                1585                1590
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Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ala Ser  Ser Ala Pro
    1595                1600                1605

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1610                1615                1620

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1625                1630                1635

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1640                1645                1650

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1655                1660                1665

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1670                1675                1680

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1685                1690                1695

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1700                1705                1710

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1715                1720                1725

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1730                1735                1740

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1745                1750                1755

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1760                1765                1770

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1775                1780                1785

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1790                1795                1800

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1805                1810                1815

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1820                1825                1830

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1835                1840                1845
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Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1850                1855                1860

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1865                1870                1875

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1880                1885                1890

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1895                1900                1905

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1910                1915                1920

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1925                1930                1935

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1940                1945                1950

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1955                1960                1965

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1970                1975                1980

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1985                1990                1995

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    2000                2005                2010

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    2015                2020                2025

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    2030                2035                2040

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    2045                2050                2055

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2060                2065                2070

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    2075                2080                2085

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    2090                2095                2100
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Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    2105                 2110                 2115

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    2120                 2125                 2130

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    2135                 2140                 2145

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2150                 2155                 2160

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2165                 2170                 2175

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    2180                 2185                 2190

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2195                 2200                 2205

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    2210                 2215                 2220

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    2225                 2230                 2235

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    2240                 2245                 2250

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    2255                 2260                 2265

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    2270                 2275                 2280

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Glu Pro Ala
    2285                 2290                 2295

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2300                 2305                 2310

Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    2315                 2320                 2325

Pro Ala  Ser Ser Glu Gln Lys  Leu Ile Ser Glu Glu  Asp Leu Ser
    2330                 2335                 2340

Pro Ala  Thr Gly His His His  His His His
    2345                 2350
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<211> 7062
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<213> Artificial Sequence

<220>
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polynucleotide

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acggggacgg gaccgggggc gtcaccagga acgtcctcca ccggctcgcc gggtgcatca 180

cccggaacgt catcgaccgg atcgccaggg agctcgacgc catcaggcgc aacaggatca 240

cctggctcaa gccctagcgc gtcaaccggc acgggtccgg gtgcctcccc tggcacgtcc 300

agcaccggat cacccggatc gagcccatcc gcctcaaccg gaaccggacc cggtacacca 360

gggtcgggaa cagcctcctc gtcaccaggc tcctcaaccc cctcgggagc cacgggttcg 420

cccggttcgt caacgccttc cggagcaact ggtagccccg gagcatcgcc aggaacttcg 480

agcacggggt cgcccgcctc gagcagtgat ctcggtgagc tgcctgtgga cgcaagattt 540

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tttgtagaat tcacggatca ccttttcaac atcgctaagc caaggccacc ctggatgggt 660

ctgctaggtc ctaccatcca ggctgaggtt tatgatacag tggtcattac acttaagaac 720

atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag 780

ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct 840

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ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

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gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg 1200

aaatcagtct attggcatgt gattggaatg ggcaccactc ctgaagtgca ctcaatattc 1260

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tcagccccag gcaccagcac ggagccctct gagggaagcg cacctggtac cagcgaaagc 2040

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<211> 2356
<212> PRT
<213> Artificial Sequence

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Met Gln Gly Ala Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
20 25 30

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
35 40 45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser

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    50                  55                  60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65                  70                  75                  80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
                85                  90                  95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
            100                 105                 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
        115                 120                 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
    130                 135                 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145                 150                 155                 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
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325 330 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
340 345 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
355 360 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
370 375 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385 390 395 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
405 410 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
420 425 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
435 440 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
450 455 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465 470 475 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
485 490 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
500 505 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
515 520 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
530 535 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu
545 550 555 560

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
565 570 575

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
580 585 590

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser

595 600 605

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
610 615 620

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
625 630 635 640

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
645 650 655

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
660 665 670

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
675 680 685

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp
690 695 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705 710 715 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
725 730 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
740 745 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
755 760 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
770 775 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785 790 795 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
805 810 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
820 825 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
835 840 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
850 855 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp

865 870 875 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
885 890 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
900 905 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
915 920 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
930 935 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945 950 955 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
965 970 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
980 985 990

His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val
995 1000 1005

Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr
1010 1015 1020

Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu
1025 1030 1035

Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His Gln
1040 1045 1050

Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
1055 1060 1065

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
1070 1075 1080

Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln
1085 1090 1095

Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp
1100 1105 1110

Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala
1115 1120 1125

Gln Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser

1130 1135 1140

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
1145 1150 1155

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
1160 1165 1170

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
1175 1180 1185

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
1190 1195 1200

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
1205 1210 1215

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
1220 1225 1230

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
1235 1240 1245

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
1250 1255 1260

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser
1265 1270 1275

Ser Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val
1280 1285 1290

Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg
1295 1300 1305

Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
1310 1315 1320

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala
1325 1330 1335

Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu
1340 1345 1350

Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro
1355 1360 1365

Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala
1370 1375 1380

Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser

1385 1390 1395

Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
1400 1405 1410

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg
1415 1420 1425

Gln Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp
1430 1435 1440

Glu Thr Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys
1445 1450 1455

Arg Gly Ala Pro Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
1460 1465 1470

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser
1475 1480 1485

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr
1490 1495 1500

Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
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Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
1520 1525 1530

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro
1535 1540 1545

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser
1550 1555 1560

Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser
1565 1570 1575

Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
1580 1585 1590

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser
1595 1600 1605

Ser Ala Pro Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu
1610 1615 1620

Asn Tyr Arg Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu
1625 1630 1635

Pro Gly Leu Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu

1640 1645 1650

Leu Ser Met Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser
1655 1660 1665

Gly His Val Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala
1670 1675 1680

Leu Tyr Asn Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu
1685 1690 1695

Pro Ser Lys Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu
1700 1705 1710

His Leu His Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn
1715 1720 1725

Lys Cys Gln Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp
1730 1735 1740

Phe Gln Ile Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys
1745 1750 1755

Leu Ala Arg Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr
1760 1765 1770

Lys Glu Pro Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met
1775 1780 1785

Ile Ile His Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser
1790 1795 1800

Ser Leu Tyr Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly
1805 1810 1815

Lys Lys Trp Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met
1820 1825 1830

Val Phe Phe Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile
1835 1840 1845

Phe Asn Pro Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr
1850 1855 1860

His Tyr Ser Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys
1865 1870 1875

Asp Leu Asn Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala
1880 1885 1890

Ile Ser Asp Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met

1895 1900 1905

Phe Ala Thr Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly
1910 1915 1920

Arg Ser Asn Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp
1925 1930 1935

Leu Gln Val Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr
1940 1945 1950

Thr Gln Gly Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu
1955 1960 1965

Phe Leu Ile Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe
1970 1975 1980

Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser
1985 1990 1995

Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg
2000 2005 2010

Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu
2015 2020 2025

Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala
2030 2035 2040

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2045 2050 2055

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
2060 2065 2070

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
2075 2080 2085

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
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Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
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Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
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Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu
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Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
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Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
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Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
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Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala
2240 2245 2250

Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
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Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2270 2275 2280

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
2285 2290 2295

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
2300 2305 2310

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
2315 2320 2325

Gly Ser Ala Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu
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Asp Leu Ser Pro Ala Thr Gly His His His His His His
2345 2350 2355

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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
35 40 45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
50 55 60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65 70 75 80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
85 90 95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
100 105 110

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Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
        115                 120                 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
    130                 135                 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145                 150                 155                 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380
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Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
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Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655
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Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
        915                 920                 925
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Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
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Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
945                 950                 955                 960

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
                965                 970                 975

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly
            980                 985                 990

Thr Ser Ser Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
        995                 1000                1005

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1010                1015                1020

Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser
    1025                1030                1035

Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
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Ser Ser  Thr Gly Ser Pro Ala  Ser Ser Ser Asp Gln  Glu Glu Ile
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Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Gly  Ala Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1130                1135                1140

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1145                1150                1155

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
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Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1175                1180                1185
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Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
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Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1220                 1225                 1230

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1235                 1240                 1245

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1250                 1255                 1260

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
    1265                 1270                 1275

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1280                 1285                 1290

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1295                 1300                 1305

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
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Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1325                 1330                 1335

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1340                 1345                 1350

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
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Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1370                 1375                 1380

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1385                 1390                 1395

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1400                 1405                 1410

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1415                 1420                 1425

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1430                 1435                 1440
```

```
Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
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Arg Gly  Ala Pro Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1460                1465                1470

Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser
    1475                1480                1485

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
    1490                1495                1500

Ser Gly  Ser Glu Thr Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1505                1510                1515

Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1520                1525                1530

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Pro
    1535                1540                1545

Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Ser Pro  Ala Gly Ser
    1550                1555                1560

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1565                1570                1575

Thr Glu  Glu Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1580                1585                1590

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Ala Ser
    1595                1600                1605

Ser Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1610                1615                1620

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1625                1630                1635

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1640                1645                1650

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1655                1660                1665

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1670                1675                1680

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1685                1690                1695
```

```
Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1700                1705                1710

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1715                1720                1725

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1730                1735                1740

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1745                1750                1755

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1760                1765                1770

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1775                1780                1785

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1790                1795                1800

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1805                1810                1815

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1820                1825                1830

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1835                1840                1845

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1850                1855                1860

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1865                1870                1875

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1880                1885                1890

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1895                1900                1905

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1910                1915                1920

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1925                1930                1935

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1940                1945                1950
```

```
Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1955                 1960                 1965

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1970                 1975                 1980

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1985                 1990                 1995

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    2000                 2005                 2010

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    2015                 2020                 2025

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    2030                 2035                 2040

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2045                 2050                 2055

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    2060                 2065                 2070

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    2075                 2080                 2085

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2090                 2095                 2100

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    2105                 2110                 2115

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    2120                 2125                 2130

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    2135                 2140                 2145

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2150                 2155                 2160

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser
    2165                 2170                 2175

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Thr Glu
    2180                 2185                 2190

Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2195                 2200                 2205
```

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2210 2215 2220

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2225 2230 2235

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala
2240 2245 2250

Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
2255 2260 2265

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2270 2275 2280

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
2285 2290 2295

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
2300 2305 2310

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
2315 2320 2325

Gly Ser Ala Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu
2330 2335 2340

Asp Leu Ser Pro Ala Thr Gly His His His His His His
2345 2350 2355

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<211> 7071
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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acggggacgg gaccgggggc gtcaccagga acgtcctcca ccggctcgcc gggtgcatca 180

cccggaacgt catcgaccgg atcgccaggg agctcgacgc catcaggcgc aacaggatca 240

cctggctcaa gccctagcgc gtcaaccggc acgggtccgg gtgcctcccc tggcacgtcc 300

agcaccggat cacccggatc gagcccatcc gcctcaaccg gaaccggacc cggtacacca 360

gggtcgggaa cagcctcctc gtcaccaggc tcctcaaccc cctcgggagc cacgggttcg 420

cccggttcgt caacgccttc cggagcaact ggtagccccg gagcatcgcc aggaacttcg 480

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<212> PRT
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Met Gln Gly Ala Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
35 40 45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
50 55 60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65 70 75 80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
85 90 95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
100 105 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
115 120 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
130 135 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145 150 155 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
165 170 175

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Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445
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Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu
545                 550                 555                 560

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
                565                 570                 575

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            580                 585                 590

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
        595                 600                 605

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    610                 615                 620

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
625                 630                 635                 640

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
                645                 650                 655

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
            660                 665                 670

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
        675                 680                 685

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp
    690                 695                 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720
```

```
Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990
```

```
His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020

Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Gly Ala Pro  Gly Thr Pro
    1055                1060                1065

Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly Ala Ser  Pro Gly Thr
    1070                1075                1080

Ser Ser  Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
    1085                1090                1095

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1100                1105                1110

Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr Gly Pro  Gly Thr Pro
    1115                1120                1125

Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly Ala Ser  Pro Gly Thr
    1130                1135                1140

Ser Ser  Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
    1145                1150                1155

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1160                1165                1170

Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser
    1175                1180                1185

Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    1190                1195                1200

Ser Ser  Thr Gly Ser Pro Ala  Ser Ser Ser Asp Gln  Glu Glu Ile
    1205                1210                1215

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1220                1225                1230

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1235                1240                1245
```

```
Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1250                1255                1260

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1265                1270                1275

Gln Gly  Ala Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1280                1285                1290

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1295                1300                1305

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1310                1315                1320

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1325                1330                1335

Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1340                1345                1350

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1355                1360                1365

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1370                1375                1380

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1385                1390                1395

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1400                1405                1410

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
    1415                1420                1425

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1430                1435                1440

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1445                1450                1455

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1460                1465                1470

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1475                1480                1485

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1490                1495                1500
```

```
Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1505                1510                1515

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1520                1525                1530

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1535                1540                1545

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1550                1555                1560

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1565                1570                1575

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1580                1585                1590

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1595                1600                1605

Arg Gly  Ala Pro Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1610                1615                1620

Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser
    1625                1630                1635

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
    1640                1645                1650

Ser Gly  Ser Glu Thr Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1655                1660                1665

Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1670                1675                1680

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Pro
    1685                1690                1695

Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Ser Pro  Ala Gly Ser
    1700                1705                1710

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1715                1720                1725

Thr Glu  Glu Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1730                1735                1740

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Ala Ser
    1745                1750                1755
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Ser Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1760                1765                1770

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1775                1780                1785

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1790                1795                1800

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1805                1810                1815

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1820                1825                1830

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1835                1840                1845

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1850                1855                1860

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1865                1870                1875

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1880                1885                1890

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1895                1900                1905

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1910                1915                1920

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1925                1930                1935

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1940                1945                1950

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1955                1960                1965

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1970                1975                1980

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1985                1990                1995

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    2000                2005                2010
```

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His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    2015                2020                2025

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    2030                2035                2040

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    2045                2050                2055

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    2060                2065                2070

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    2075                2080                2085

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    2090                2095                2100

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    2105                2110                2115

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    2120                2125                2130

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    2135                2140                2145

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    2150                2155                2160

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    2165                2170                2175

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    2180                2185                2190

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2195                2200                2205

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    2210                2215                2220

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    2225                2230                2235

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2240                2245                2250

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    2255                2260                2265
```

```
Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    2270                 2275                 2280

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    2285                 2290                 2295

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2300                 2305                 2310

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser
    2315                 2320                 2325

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Thr Glu
    2330                 2335                 2340

Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2345                 2350                 2355

Glu Ser  Gly Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2360                 2365                 2370

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2375                 2380                 2385

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Ser  Glu Pro Ala
    2390                 2395                 2400

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Pro Ala Gly  Ser Pro Thr
    2405                 2410                 2415

Ser Thr  Glu Glu Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    2420                 2425                 2430

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    2435                 2440                 2445

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    2450                 2455                 2460

Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu
    2465                 2470                 2475

Gly Ser  Ala Pro Ala Ser Ser  Glu Gln Lys Leu Ile  Ser Glu Glu
    2480                 2485                 2490

Asp Leu  Ser Pro Ala Thr Gly  His His His His His  His
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<213> Artificial Sequence

<220>
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<211> 2356
<212> PRT
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        35                  40                  45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
    50                  55                  60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65                  70                  75                  80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
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Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
            100                 105                 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
        115                 120                 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
    130                 135                 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145                 150                 155                 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285
```

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Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu
545                 550                 555                 560
```

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Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
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Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            580                 585                 590

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
        595                 600                 605

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    610                 615                 620

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
625                 630                 635                 640

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
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Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
            660                 665                 670

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
        675                 680                 685

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp
    690                 695                 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830
```

```
Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
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Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020

Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Gly Ala Pro Ser Pro  Ala Gly Ser
    1040                1045                1050

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1055                1060                1065

Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro
    1070                1075                1080

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1085                1090                1095
```

```
Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Thr Glu Pro
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Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
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Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
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Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
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Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
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Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1175                1180                1185

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1190                1195                1200

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1205                1210                1215

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1220                1225                1230

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1235                1240                1245

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1250                1255                1260

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1265                1270                1275

Gln Gly  Ala Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1280                1285                1290

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1295                1300                1305

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
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Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
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Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
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Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
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Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
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Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1385                 1390                 1395

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
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Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
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Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
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Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
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Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
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Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
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Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
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Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
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Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
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Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1535                 1540                 1545

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1550                 1555                 1560

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1565                 1570                 1575

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1580                 1585                 1590

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1595                 1600                 1605
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Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
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Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
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Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1640                 1645                 1650

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1655                 1660                 1665

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1670                 1675                 1680

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1685                 1690                 1695

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1700                 1705                 1710

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1715                 1720                 1725

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1730                 1735                 1740

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1745                 1750                 1755

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1760                 1765                 1770

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1775                 1780                 1785

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1790                 1795                 1800

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1805                 1810                 1815

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1820                 1825                 1830

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
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Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
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His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
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Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
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Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1895                1900                1905

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1910                1915                1920

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1925                1930                1935

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1940                1945                1950

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1955                1960                1965

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1970                1975                1980

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1985                1990                1995

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    2000                2005                2010

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    2015                2020                2025

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
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Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
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Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
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Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2090                2095                2100

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
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Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    2120                 2125                 2130

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
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Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2150                 2155                 2160

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser
    2165                 2170                 2175

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Thr Glu
    2180                 2185                 2190

Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2195                 2200                 2205

Glu Ser  Gly Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2210                 2215                 2220

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2225                 2230                 2235

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Ser  Glu Pro Ala
    2240                 2245                 2250

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Pro Ala Gly  Ser Pro Thr
    2255                 2260                 2265

Ser Thr  Glu Glu Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    2270                 2275                 2280

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
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Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
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Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu
    2315                 2320                 2325

Gly Ser  Ala Pro Ala Ser Ser  Glu Gln Lys Leu Ile  Ser Glu Glu
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Asp Leu  Ser Pro Ala Thr Gly  His His His His His  His
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<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1173

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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
        35                  40                  45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
    50                  55                  60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
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65 70 75 80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
85 90 95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
100 105 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
115 120 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
130 135 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145 150 155 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
260 265 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
275 280 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
290 295 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305 310 315 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
325 330 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe

340 345 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
355 360 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
370 375 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385 390 395 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
405 410 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
420 425 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
435 440 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
450 455 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465 470 475 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
485 490 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
500 505 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
515 520 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
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Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu
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Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
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Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
580 585 590

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595 600 605

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly

610 615 620

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
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Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
645 650 655

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
660 665 670

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
675 680 685

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp
690 695 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705 710 715 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
725 730 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
740 745 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
755 760 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
770 775 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785 790 795 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
805 810 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
820 825 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
835 840 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
850 855 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865 870 875 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro

885 890 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
900 905 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
915 920 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
930 935 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945 950 955 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
965 970 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
980 985 990

His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val
995 1000 1005

Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr
1010 1015 1020

Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu
1025 1030 1035

Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly Ser
1040 1045 1050

Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
1055 1060 1065

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
1070 1075 1080

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
1085 1090 1095

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro
1100 1105 1110

Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
1115 1120 1125

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
1130 1135 1140

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser

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Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1175                1180                1185

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1190                1195                1200

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1205                1210                1215

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1220                1225                1230

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1235                1240                1245

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1250                1255                1260

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1265                1270                1275

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1280                1285                1290

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1295                1300                1305

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1310                1315                1320

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1325                1330                1335

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1340                1345                1350

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1355                1360                1365

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1370                1375                1380

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1385                1390                1395

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
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1400 1405 1410

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1415 1420 1425

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1430 1435 1440

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Gly Ala
1445 1450 1455

Pro Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
1460 1465 1470

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala
1475 1480 1485

Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
1490 1495 1500

Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
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Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
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Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser
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Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser
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Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
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Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Ala Pro
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Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1610 1615 1620

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1625 1630 1635

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1640 1645 1650

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val

1655 1660 1665

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1670 1675 1680

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1685 1690 1695

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1700 1705 1710

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1715 1720 1725

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1730 1735 1740

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1745 1750 1755

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1760 1765 1770

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1775 1780 1785

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1790 1795 1800

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1805 1810 1815

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1820 1825 1830

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1835 1840 1845

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1850 1855 1860

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1865 1870 1875

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1880 1885 1890

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1895 1900 1905

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn

1910 1915 1920

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1925 1930 1935

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1940 1945 1950

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1955 1960 1965

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1970 1975 1980

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1985 1990 1995

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
2000 2005 2010

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
2015 2020 2025

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Pro Gly Thr
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2060 2065 2070

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
2075 2080 2085

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
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2105 2110 2115

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
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Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
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Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
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Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly

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Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
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Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
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Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
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Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
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Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
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Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Ala Thr Gly His His His His His His
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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
35 40 45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
50 55 60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65 70 75 80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
85 90 95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
100 105 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
115 120 125

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Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
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Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145                 150                 155                 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400
```

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Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
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His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670
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Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940
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Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
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Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200
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Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
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Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                 1225                 1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                 1240                 1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                 1255                 1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                 1270                 1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                 1285                 1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1295                 1300                 1305

Pro Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    1310                 1315                 1320

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1325                 1330                 1335

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1340                 1345                 1350

Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1355                 1360                 1365

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1370                 1375                 1380

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Pro  Ala Gly Ser
    1385                 1390                 1395

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1400                 1405                 1410

Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1415                 1420                 1425

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1430                 1435                 1440

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ala Ser  Ser Ala Pro
    1445                 1450                 1455
```

```
Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
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Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                 1480                 1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                 1495                 1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                 1510                 1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                 1525                 1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                 1540                 1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                 1555                 1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                 1570                 1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                 1585                 1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                 1600                 1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                 1615                 1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                 1630                 1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                 1645                 1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                 1660                 1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                 1675                 1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                 1690                 1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
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```
Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
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Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                1960                1965
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Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
1970 1975 1980

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
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Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2000 2005 2010

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
2015 2020 2025

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
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Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
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Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
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Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
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Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
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Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
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Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
35 40 45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
50 55 60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65 70 75 80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
85 90 95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
100 105 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
115 120 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser

130 135 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145 150 155 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
260 265 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
275 280 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
290 295 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305 310 315 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
325 330 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
340 345 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
355 360 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
370 375 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385 390 395 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val

405 410 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
420 425 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
435 440 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
450 455 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465 470 475 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
485 490 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
500 505 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
515 520 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
530 535 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545 550 555 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
565 570 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
580 585 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
595 600 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
610 615 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
645 650 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn

675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
835 840 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
885 890 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
900 905 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
915 920 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
930 935 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser

945 950 955 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
965 970 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
980 985 990

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
995 1000 1005

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser
1010 1015 1020

Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
1025 1030 1035

Ser Ala Pro Gly Ala Ser Ser Pro Pro Val Leu Lys Arg His Gln
1040 1045 1050

Ala Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
1055 1060 1065

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
1070 1075 1080

Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln
1085 1090 1095

Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp
1100 1105 1110

Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala
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Gln Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
1130 1135 1140

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
1145 1150 1155

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
1160 1165 1170

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
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Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
1190 1195 1200

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala

1205 1210 1215

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
1220 1225 1230

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
1235 1240 1245

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
1250 1255 1260

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser
1265 1270 1275

Ser Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val
1280 1285 1290

Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg
1295 1300 1305

Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
1310 1315 1320

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala
1325 1330 1335

Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu
1340 1345 1350

Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro
1355 1360 1365

Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala
1370 1375 1380

Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser
1385 1390 1395

Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
1400 1405 1410

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg
1415 1420 1425

Gln Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp
1430 1435 1440

Glu Thr Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys
1445 1450 1455

Arg Gly Ala Pro Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro

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Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser
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    1490                1495                1500

Ser Gly  Ser Glu Thr Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
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Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
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Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Pro
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Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Ser Pro  Ala Gly Ser
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Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
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Thr Glu  Glu Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1580                1585                1590

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Ala Ser
    1595                1600                1605

Ser Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1610                1615                1620

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
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Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1640                1645                1650

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1655                1660                1665

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1670                1675                1680

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1685                1690                1695

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1700                1705                1710

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
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Lys Cys Gln Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp
1730 1735 1740

Phe Gln Ile Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys
1745 1750 1755

Leu Ala Arg Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr
1760 1765 1770

Lys Glu Pro Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met
1775 1780 1785

Ile Ile His Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser
1790 1795 1800

Ser Leu Tyr Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly
1805 1810 1815

Lys Lys Trp Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met
1820 1825 1830

Val Phe Phe Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile
1835 1840 1845

Phe Asn Pro Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr
1850 1855 1860

His Tyr Ser Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys
1865 1870 1875

Asp Leu Asn Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala
1880 1885 1890

Ile Ser Asp Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met
1895 1900 1905

Phe Ala Thr Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly
1910 1915 1920

Arg Ser Asn Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp
1925 1930 1935

Leu Gln Val Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr
1940 1945 1950

Thr Gln Gly Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu
1955 1960 1965

Phe Leu Ile Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe

1970 1975 1980

Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser
1985 1990 1995

Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg
2000 2005 2010

Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu
2015 2020 2025

Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala
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Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
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Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
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Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
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Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
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Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
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Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
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Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser
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Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
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Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser

2225 2230 2235

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala
2240 2245 2250

Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
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Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2270 2275 2280

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
2285 2290 2295

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
2300 2305 2310

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
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Gly Ser Ala Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu
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Asp Leu Ser Pro Ala Thr Gly His His His His His His
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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
35 40 45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
50 55 60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65 70 75 80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
85 90 95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
100 105 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
115 120 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
130 135 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145 150 155 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

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Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460
```

```
His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu
545                 550                 555                 560

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
                565                 570                 575

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            580                 585                 590

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
        595                 600                 605

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    610                 615                 620

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
625                 630                 635                 640

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
                645                 650                 655

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
            660                 665                 670

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
        675                 680                 685

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp
    690                 695                 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735
```

```
Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005
```

```
Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020

Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Gly Ala Pro Ser Pro  Ala Gly Ser
    1040                1045                1050

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1055                1060                1065

Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro
    1070                1075                1080

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1085                1090                1095

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Thr Glu Pro
    1100                1105                1110

Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1115                1120                1125

Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1130                1135                1140

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1145                1150                1155

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1160                1165                1170

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1175                1180                1185

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1190                1195                1200

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1205                1210                1215

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1220                1225                1230

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1235                1240                1245

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1250                1255                1260
```

```
Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1265                1270                1275

Gln Gly  Ala Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1280                1285                1290

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1295                1300                1305

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1310                1315                1320

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1325                1330                1335

Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1340                1345                1350

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1355                1360                1365

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1370                1375                1380

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1385                1390                1395

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1400                1405                1410

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
    1415                1420                1425

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1430                1435                1440

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1445                1450                1455

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1460                1465                1470

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1475                1480                1485

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1490                1495                1500

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1505                1510                1515
```

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Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1520                1525                1530

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1535                1540                1545

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1550                1555                1560

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1565                1570                1575

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1580                1585                1590

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1595                1600                1605

Arg Gly  Ala Pro Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1610                1615                1620

Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser
    1625                1630                1635

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
    1640                1645                1650

Ser Gly  Ser Glu Thr Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1655                1660                1665

Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1670                1675                1680

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Pro
    1685                1690                1695

Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Ser Pro  Ala Gly Ser
    1700                1705                1710

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1715                1720                1725

Thr Glu  Glu Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1730                1735                1740

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Ala Ser
    1745                1750                1755

Ser Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1760                1765                1770
```

```
Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1775                1780                1785

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1790                1795                1800

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1805                1810                1815

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1820                1825                1830

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1835                1840                1845

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1850                1855                1860

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1865                1870                1875

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1880                1885                1890

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1895                1900                1905

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1910                1915                1920

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1925                1930                1935

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1940                1945                1950

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1955                1960                1965

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1970                1975                1980

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1985                1990                1995

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    2000                2005                2010

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    2015                2020                2025
```

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Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    2030                2035                2040

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    2045                2050                2055

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    2060                2065                2070

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    2075                2080                2085

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    2090                2095                2100

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    2105                2110                2115

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    2120                2125                2130

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    2135                2140                2145

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    2150                2155                2160

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    2165                2170                2175

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    2180                2185                2190

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
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Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    2210                2215                2220

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    2225                2230                2235

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
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Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    2255                2260                2265

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    2270                2275                2280
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Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
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Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser
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Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu
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Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
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Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala
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Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
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Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2420 2425 2430

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
2435 2440 2445

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
2450 2455 2460

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300
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Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
                405                 410                 415

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            420                 425                 430

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
        435                 440                 445

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    450                 455                 460

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
465                 470                 475                 480

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
                485                 490                 495

Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            500                 505                 510

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
        515                 520                 525

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
    530                 535                 540

Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575
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Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845
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Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110
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Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1295                1300                1305

Pro Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    1310                1315                1320

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1325                1330                1335

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
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Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1355                1360                1365
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Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
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Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Pro  Ala Gly Ser
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Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1400                 1405                 1410

Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1415                 1420                 1425

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1430                 1435                 1440

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ala Ser  Ser Ala Pro
    1445                 1450                 1455

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                 1465                 1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                 1480                 1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                 1495                 1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                 1510                 1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                 1525                 1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                 1540                 1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                 1555                 1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                 1570                 1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                 1585                 1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                 1600                 1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                 1615                 1620
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Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
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Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875
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Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                 1885                 1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                 1900                 1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                 1915                 1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                 1930                 1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                 1945                 1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                 1960                 1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                 1975                 1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                 1990                 1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                 2005                 2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                 2020                 2025

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    2030                 2035                 2040

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2045                 2050                 2055

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    2060                 2065                 2070

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    2075                 2080                 2085

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    2090                 2095                 2100

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    2105                 2110                 2115

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    2120                 2125                 2130
```

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
2135 2140 2145

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2150 2155 2160

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2165 2170 2175

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
2180 2185 2190

Pro Ala Thr Gly His His His His His His
2195 2200

<210> 1182
<211> 6612
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg gcgcgccaac cagtacggag ccgtccgagg ggagcgcacc aggaagcccg 1260

gctgggagcc cgacttctac cgaagagggt acatctaccg aaccaagtga aggttcagca 1320

ccaggcacct caacagaacc ctctgagggc tcggcgcctg gtacaagtga gtccgccacc 1380

ccagaatccg ggcctgggac aagcacagaa ccttcggaag ggagtgcccc tggaacatcc 1440

gaatcggcaa ccccagaatc agggccagga tctgagcccg cgacttcggg ctccgagacg 1500

cctgggacat ccaccgagcc ctccgaagga tcagccccag gcaccagcac ggagccctct 1560

gagggaagcg cacctggtac cagcgaaagc gcaactcccg aatcaggtcc cggtacgagc 1620

gagtcggcga ccccggagag cgggccaggt gcctcgagcg acagaagtta taaaagtcaa 1680

tatttgaaca atggccctca gcggattggt aggaagtaca aaaaagtccg atttatggca 1740

tacacagatg aaacctttaa gactcgtgaa gctattcagc atgaatcagg aatcttggga 1800

cctttacttt atggggaagt tggagacaca ctgttgatta tatttaagaa tcaagcaagc 1860

agaccatata acatctaccc tcacggaatc actgatgtcc gtcctttgta ttcaaggaga 1920

ttaccaaaag gtgtaaaaca tttgaaggat tttccaattc tgccaggaga aatattcaaa 1980

tataaatgga cagtgactgt agaagatggg ccaactaaat cagatcctcg gtgcctgacc 2040

cgctattact ctagtttcgt taatatggag agagatctag cttcaggact cattggccct 2100

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catttgggac tcctggggcc atatataaga gcagaagttg aagataatat catggtaact 3540

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gatcagaggc aaggagcaga acctagaaaa aactttgtca agcctaatga aaccaaaact 3660

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ccatccgagg gatcagctcc cggcacctca gagtcggcaa ccccggagtc tggacccgga 6180

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gccacccccg agagtggccc aggcactagt accgagccct ctgaaggcag tgcgccagcc 6540

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catcaccatt ga 6612

<210> 1183
<211> 2356
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1183
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

```
Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300
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```
Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
                405                 410                 415

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            420                 425                 430

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
        435                 440                 445

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    450                 455                 460

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
465                 470                 475                 480

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
                485                 490                 495

Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            500                 505                 510

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
        515                 520                 525

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
    530                 535                 540

Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575
```

```
Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845
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Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110
```

```
Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Gly  Ala Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1130                1135                1140

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1145                1150                1155

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1160                1165                1170

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1175                1180                1185

Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1190                1195                1200

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1205                1210                1215

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1220                1225                1230

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1235                1240                1245

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1250                1255                1260

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
    1265                1270                1275

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1280                1285                1290

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1295                1300                1305

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1310                1315                1320

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1325                1330                1335

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1340                1345                1350

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1355                1360                1365
```

```
Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1370                1375                1380

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1385                1390                1395

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1400                1405                1410

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1415                1420                1425

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1430                1435                1440

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1445                1450                1455

Arg Gly  Ala Pro Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1460                1465                1470

Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser
    1475                1480                1485

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
    1490                1495                1500

Ser Gly  Ser Glu Thr Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1505                1510                1515

Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1520                1525                1530

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Pro
    1535                1540                1545

Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Ser Pro  Ala Gly Ser
    1550                1555                1560

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1565                1570                1575

Thr Glu  Glu Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1580                1585                1590

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Ala Ser
    1595                1600                1605

Ser Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1610                1615                1620
```

```
Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1625                1630                1635

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1640                1645                1650

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1655                1660                1665

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1670                1675                1680

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1685                1690                1695

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1700                1705                1710

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1715                1720                1725

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1730                1735                1740

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1745                1750                1755

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1760                1765                1770

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1775                1780                1785

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1790                1795                1800

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1805                1810                1815

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1820                1825                1830

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1835                1840                1845

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1850                1855                1860

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1865                1870                1875
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```
Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1880                1885                1890

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1895                1900                1905

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1910                1915                1920

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1925                1930                1935

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1940                1945                1950

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1955                1960                1965

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1970                1975                1980

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1985                1990                1995

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    2000                2005                2010

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    2015                2020                2025

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    2030                2035                2040

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2045                2050                2055

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    2060                2065                2070

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    2075                2080                2085

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2090                2095                2100

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    2105                2110                2115

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    2120                2125                2130
```

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
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Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
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Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser
2165 2170 2175

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu
2180 2185 2190

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
2195 2200 2205

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2210 2215 2220

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2225 2230 2235

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala
2240 2245 2250

Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
2255 2260 2265

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2270 2275 2280

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
2285 2290 2295

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
2300 2305 2310

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
2315 2320 2325

Gly Ser Ala Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu
2330 2335 2340

Asp Leu Ser Pro Ala Thr Gly His His His His His His
2345 2350 2355

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<211> 7071
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<213> Artificial Sequence

<220>
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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

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ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

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20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

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Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365
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Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640
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Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
            740                 745                 750

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
        755                 760                 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
    770                 775                 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785                 790                 795                 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                805                 810                 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            820                 825                 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
        835                 840                 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    850                 855                 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865                 870                 875                 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910
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Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
            980                 985                 990

Ser Pro Gly Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
        995                 1000                1005

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1010                1015                1020

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1025                1030                1035

Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1040                1045                1050

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1055                1060                1065

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1070                1075                1080

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1085                1090                1095

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1100                1105                1110

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
    1115                1120                1125

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1130                1135                1140

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1145                1150                1155

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1160                1165                1170
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Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1175                1180                1185

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1190                1195                1200

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1205                1210                1215

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1220                1225                1230

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1235                1240                1245

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1250                1255                1260

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1265                1270                1275

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1280                1285                1290

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1295                1300                1305

Arg Gly  Ala Pro Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1310                1315                1320

Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser
    1325                1330                1335

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
    1340                1345                1350

Ser Gly  Ser Glu Thr Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1355                1360                1365

Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1370                1375                1380

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Pro
    1385                1390                1395

Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Ser Pro  Ala Gly Ser
    1400                1405                1410

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1415                1420                1425
```

```
Thr Glu  Glu Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1430                1435                1440

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Ala Ser
    1445                1450                1455

Ser Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1460                1465                1470

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1475                1480                1485

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1490                1495                1500

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1505                1510                1515

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1520                1525                1530

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1535                1540                1545

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1550                1555                1560

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1565                1570                1575

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1580                1585                1590

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1595                1600                1605

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1610                1615                1620

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1625                1630                1635

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1640                1645                1650

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1655                1660                1665

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1670                1675                1680
```

```
Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1685                1690                1695

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1700                1705                1710

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1715                1720                1725

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1730                1735                1740

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1745                1750                1755

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1760                1765                1770

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1775                1780                1785

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1790                1795                1800

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1805                1810                1815

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1820                1825                1830

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1835                1840                1845

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    1850                1855                1860

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    1865                1870                1875

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    1880                1885                1890

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1895                1900                1905

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    1910                1915                1920

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1925                1930                1935
```

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Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1940                1945                1950

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    1955                1960                1965

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    1970                1975                1980

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1985                1990                1995

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2000                2005                2010

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser
    2015                2020                2025

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Thr Glu
    2030                2035                2040

Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2045                2050                2055

Glu Ser  Gly Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2060                2065                2070

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2075                2080                2085

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Ser  Glu Pro Ala
    2090                2095                2100

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Pro Ala Gly  Ser Pro Thr
    2105                2110                2115

Ser Thr  Glu Glu Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    2120                2125                2130

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    2135                2140                2145

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    2150                2155                2160

Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu
    2165                2170                2175

Gly Ser  Ala Pro Ala Ser Ser  Glu Gln Lys Leu Ile  Ser Glu Glu
    2180                2185                2190
```

Asp Leu Ser Pro Ala Thr Gly His His His His His His
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<210> 1186
<211> 6621
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

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atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

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gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

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acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

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<210> 1187
<211> 2203
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Gly Ser Ser
20 25 30

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
35 40 45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
50 55 60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65 70 75 80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
85 90 95

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Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
            100                 105                 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
        115                 120                 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
    130                 135                 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145                 150                 155                 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365
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Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu
545                 550                 555                 560

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
                565                 570                 575

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            580                 585                 590

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
        595                 600                 605

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    610                 615                 620

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
625                 630                 635                 640
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Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
                645                 650                 655

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
            660                 665                 670

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
        675                 680                 685

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp
    690                 695                 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910
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Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020

Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Gly Ala Pro  Gly Thr Pro
    1055                1060                1065

Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly Ala Ser  Pro Gly Thr
    1070                1075                1080

Ser Ser  Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
    1085                1090                1095

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1100                1105                1110

Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr Gly Pro  Gly Thr Pro
    1115                1120                1125

Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly Ala Ser  Pro Gly Thr
    1130                1135                1140

Ser Ser  Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
    1145                1150                1155

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1160                1165                1170
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Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser
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Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
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Ser Ser  Thr Gly Ser Pro Ala  Ser Ser Ser Asp Gln  Glu Glu Ile
    1205                1210                1215

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1220                1225                1230

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1235                1240                1245

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1250                1255                1260

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1265                1270                1275

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1280                1285                1290

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1295                1300                1305

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1310                1315                1320

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1325                1330                1335

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1340                1345                1350

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1355                1360                1365

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1370                1375                1380

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1385                1390                1395

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1400                1405                1410

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1415                1420                1425
```

```
Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
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Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1445                1450                1455

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                1555                1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                1615                1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680
```

```
Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
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Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                 1705                 1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                 1720                 1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                 1735                 1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                 1750                 1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                 1765                 1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                 1780                 1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                 1795                 1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                 1810                 1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                 1825                 1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                 1840                 1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                 1855                 1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                 1870                 1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                 1885                 1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                 1900                 1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                 1915                 1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                 1930                 1935
```

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1940 1945 1950

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
1955 1960 1965

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
1970 1975 1980

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1985 1990 1995

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2000 2005 2010

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
2015 2020 2025

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
2030 2035 2040

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2045 2050 2055

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
2060 2065 2070

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
2075 2080 2085

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
2090 2095 2100

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
2105 2110 2115

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
2120 2125 2130

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
2135 2140 2145

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2150 2155 2160

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2165 2170 2175

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
2180 2185 2190

Pro Ala Thr Gly His His His His His His
2195 2200

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<211> 6612
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<213> Artificial Sequence

<220>
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accggaaccg gacccggtac accagggtcg ggaacagcct cctcgtcacc aggctcctca 420

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catcaccatt ga 6612

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<212> PRT
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Gly Ser Ser
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Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
35 40 45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
50 55 60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65 70 75 80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
85 90 95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser

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            100                 105                 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
        115                 120                 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
    130                 135                 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145                 150                 155                 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
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Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
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Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
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His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
420 425 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
435 440 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
450 455 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465 470 475 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
485 490 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
500 505 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
515 520 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
530 535 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu
545 550 555 560

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
565 570 575

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
580 585 590

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595 600 605

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
610 615 620

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
625 630 635 640

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu

645 650 655

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
660 665 670

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
675 680 685

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp
690 695 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705 710 715 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
725 730 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
740 745 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
755 760 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
770 775 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785 790 795 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
805 810 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
820 825 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
835 840 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
850 855 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865 870 875 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
885 890 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
900 905 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser

915 920 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
930 935 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945 950 955 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
965 970 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
980 985 990

His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val
995 1000 1005

Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr
1010 1015 1020

Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu
1025 1030 1035

Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His Gln
1040 1045 1050

Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
1055 1060 1065

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
1070 1075 1080

Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln
1085 1090 1095

Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp
1100 1105 1110

Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala
1115 1120 1125

Gln Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
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Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
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Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
1160 1165 1170

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr

1175 1180 1185

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
1190 1195 1200

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
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Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
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Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
1235 1240 1245

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
1250 1255 1260

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser
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Ser Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val
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Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg
1295 1300 1305

Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
1310 1315 1320

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala
1325 1330 1335

Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu
1340 1345 1350

Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro
1355 1360 1365

Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala
1370 1375 1380

Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser
1385 1390 1395

Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
1400 1405 1410

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg
1415 1420 1425

Gln Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp

1430 1435 1440

Glu Thr Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys
1445 1450 1455

Arg Ala Pro Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu
1460 1465 1470

Asn Tyr Arg Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu
1475 1480 1485

Pro Gly Leu Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu
1490 1495 1500

Leu Ser Met Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser
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Gly His Val Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala
1520 1525 1530

Leu Tyr Asn Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu
1535 1540 1545

Pro Ser Lys Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu
1550 1555 1560

His Leu His Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn
1565 1570 1575

Lys Cys Gln Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp
1580 1585 1590

Phe Gln Ile Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys
1595 1600 1605

Leu Ala Arg Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr
1610 1615 1620

Lys Glu Pro Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met
1625 1630 1635

Ile Ile His Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser
1640 1645 1650

Ser Leu Tyr Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly
1655 1660 1665

Lys Lys Trp Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met
1670 1675 1680

Val Phe Phe Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile

1685 1690 1695

Phe Asn Pro Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr
1700 1705 1710

His Tyr Ser Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys
1715 1720 1725

Asp Leu Asn Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala
1730 1735 1740

Ile Ser Asp Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met
1745 1750 1755

Phe Ala Thr Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly
1760 1765 1770

Arg Ser Asn Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp
1775 1780 1785

Leu Gln Val Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr
1790 1795 1800

Thr Gln Gly Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu
1805 1810 1815

Phe Leu Ile Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe
1820 1825 1830

Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser
1835 1840 1845

Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg
1850 1855 1860

Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu
1865 1870 1875

Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala
1880 1885 1890

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
1895 1900 1905

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
1910 1915 1920

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
1925 1930 1935

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly

1940 1945 1950

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
1955 1960 1965

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
1970 1975 1980

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
1985 1990 1995

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2000 2005 2010

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser
2015 2020 2025

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu
2030 2035 2040

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
2045 2050 2055

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2060 2065 2070

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2075 2080 2085

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala
2090 2095 2100

Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
2105 2110 2115

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2120 2125 2130

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
2135 2140 2145

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
2150 2155 2160

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
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Gly Ser Ala Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu
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Asp Leu Ser Pro Ala Thr Gly His His His His His His

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tccaccggct cgccgggtgc atcacccgga acgtcatcga ccggatcgcc agggagctcg 240

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ccgggtgcct cccctggcac gtccagcacc ggatcacccg gatcgagccc atccgcctca 360

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catcatcacc atcaccattg a 6621

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<211> 2203
<212> PRT
<213> Artificial Sequence

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Gly Ser Ser
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Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
35 40 45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
50 55 60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65 70 75 80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
85 90 95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
100 105 110

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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
        115                 120                 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
    130                 135                 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145                 150                 155                 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380
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Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu
545                 550                 555                 560

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
                565                 570                 575

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            580                 585                 590

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
        595                 600                 605

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    610                 615                 620

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
625                 630                 635                 640

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
                645                 650                 655
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Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
            660                 665                 670

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
        675                 680                 685

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp
    690                 695                 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925
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Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020

Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185
```

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Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
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Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                 1210                 1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                 1225                 1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                 1240                 1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                 1255                 1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                 1270                 1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                 1285                 1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1295                 1300                 1305

Pro Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    1310                 1315                 1320

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1325                 1330                 1335

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1340                 1345                 1350

Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1355                 1360                 1365

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1370                 1375                 1380

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Pro  Ala Gly Ser
    1385                 1390                 1395

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1400                 1405                 1410

Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1415                 1420                 1425

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1430                 1435                 1440
```

```
Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ala Ser  Ser Ala Pro
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Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                1555                1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                1615                1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695
```

```
Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950
```

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Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                 1960                 1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                 1975                 1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                 1990                 1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                 2005                 2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                 2020                 2025

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    2030                 2035                 2040

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2045                 2050                 2055

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    2060                 2065                 2070

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    2075                 2080                 2085

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    2090                 2095                 2100

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    2105                 2110                 2115

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    2120                 2125                 2130

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Glu Pro Ala
    2135                 2140                 2145

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2150                 2155                 2160

Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    2165                 2170                 2175

Pro Ala  Ser Ser Glu Gln Lys  Leu Ile Ser Glu Glu  Asp Leu Ser
    2180                 2185                 2190

Pro Ala  Thr Gly His His His  His His His
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polynucleotide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

```
Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380
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Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
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Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
                405                 410                 415

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            420                 425                 430

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
        435                 440                 445

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    450                 455                 460

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
465                 470                 475                 480

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
                485                 490                 495

Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            500                 505                 510

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
        515                 520                 525

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
    530                 535                 540

Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655
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Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
        915                 920                 925
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Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
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Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
945                 950                 955                 960

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
                965                 970                 975

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly
            980                 985                 990

Thr Ser Ser Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
        995                 1000                1005

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1010                1015                1020

Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser
    1025                1030                1035

Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    1040                1045                1050

Ser Ser  Thr Gly Ser Pro Ala  Ser Ser Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Gly  Ala Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1130                1135                1140

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1145                1150                1155

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1160                1165                1170

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1175                1180                1185
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Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
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Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
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Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
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Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1235                 1240                 1245

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1250                 1255                 1260

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
    1265                 1270                 1275

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1280                 1285                 1290

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1295                 1300                 1305

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1310                 1315                 1320

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1325                 1330                 1335

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1340                 1345                 1350

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1355                 1360                 1365

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1370                 1375                 1380

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1385                 1390                 1395

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1400                 1405                 1410

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1415                 1420                 1425

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1430                 1435                 1440
```

```
Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
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Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1460                1465                1470

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1475                1480                1485

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1490                1495                1500

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1505                1510                1515

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1520                1525                1530

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1535                1540                1545

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1550                1555                1560

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1565                1570                1575

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1580                1585                1590

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1595                1600                1605

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1610                1615                1620

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1625                1630                1635

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1640                1645                1650

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1655                1660                1665

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1670                1675                1680

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1685                1690                1695
```

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Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
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His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1715                1720                1725

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1730                1735                1740

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1745                1750                1755

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1760                1765                1770

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1775                1780                1785

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1790                1795                1800

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1805                1810                1815

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1820                1825                1830

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1835                1840                1845

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    1850                1855                1860

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    1865                1870                1875

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    1880                1885                1890

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1895                1900                1905

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    1910                1915                1920

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1925                1930                1935

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1940                1945                1950
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Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    1955                 1960                 1965

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    1970                 1975                 1980

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1985                 1990                 1995

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2000                 2005                 2010

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser
    2015                 2020                 2025

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Thr Glu
    2030                 2035                 2040

Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2045                 2050                 2055

Glu Ser  Gly Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2060                 2065                 2070

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2075                 2080                 2085

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Ser  Glu Pro Ala
    2090                 2095                 2100

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Pro Ala Gly  Ser Pro Thr
    2105                 2110                 2115

Ser Thr  Glu Glu Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    2120                 2125                 2130

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    2135                 2140                 2145

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    2150                 2155                 2160

Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu
    2165                 2170                 2175

Gly Ser  Ala Pro Ala Ser Ser  Glu Gln Lys Leu Ile  Ser Glu Glu
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Asp Leu  Ser Pro Ala Thr Gly  His His His His His  His
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20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

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Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380
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        435                 440                 445

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    450                 455                 460

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
465                 470                 475                 480

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
                485                 490                 495

Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            500                 505                 510

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
        515                 520                 525

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
    530                 535                 540

Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655
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Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
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Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
        915                 920                 925
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Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
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Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
945                 950                 955                 960

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
                965                 970                 975

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly
            980                 985                 990

Thr Ser Ser Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
        995                 1000                1005

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1010                1015                1020

Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser
    1025                1030                1035

Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    1040                1045                1050

Ser Ser  Thr Gly Ser Pro Ala  Ser Ser Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185
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Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
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Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
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Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                 1225                 1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                 1240                 1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                 1255                 1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                 1270                 1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                 1285                 1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1295                 1300                 1305

Pro Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    1310                 1315                 1320

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1325                 1330                 1335

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
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Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1355                 1360                 1365

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1370                 1375                 1380

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Pro  Ala Gly Ser
    1385                 1390                 1395

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1400                 1405                 1410

Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1415                 1420                 1425

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1430                 1435                 1440
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Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ala Ser  Ser Ala Pro
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Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                1555                1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                1615                1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
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Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695
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Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
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Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
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Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
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Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950
```

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Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                 1960                 1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                 1975                 1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                 1990                 1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                 2005                 2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                 2020                 2025

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    2030                 2035                 2040

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2045                 2050                 2055

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    2060                 2065                 2070

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    2075                 2080                 2085

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    2090                 2095                 2100

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    2105                 2110                 2115

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    2120                 2125                 2130

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Glu Pro Ala
    2135                 2140                 2145

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2150                 2155                 2160

Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    2165                 2170                 2175

Pro Ala  Ser Ser Glu Gln Lys  Leu Ile Ser Glu Glu  Asp Leu Ser
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Pro Ala  Thr Gly His His His  His His His
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35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val

115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
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Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
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Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
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Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
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Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
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Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu

385 390 395 400

Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
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Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
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Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
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Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
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Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
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Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
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Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
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Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
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Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
565 570 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
580 585 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
595 600 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
610 615 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
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Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr

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Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
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Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
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Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
885 890 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
900 905 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
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Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro

930 935 940

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Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
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Arg Ala Gln Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
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Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
995 1000 1005

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
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Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser
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Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg
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Gln Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp
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Glu Thr Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys
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Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser
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Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser
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Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser

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Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
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Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
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Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
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Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
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Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
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His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
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Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
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Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
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Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
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Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
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Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
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Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
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Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
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Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
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Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
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Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
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Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
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Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
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Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
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Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
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Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
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Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    1850                1855                1860

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    1865                1870                1875

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    1880                1885                1890

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
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Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    1910                1915                1920

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1925                1930                1935

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1940                1945                1950

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
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Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
1985 1990 1995

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2000 2005 2010

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser
2015 2020 2025

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu
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Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
2045 2050 2055

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2060 2065 2070

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
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Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala
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Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
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Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
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Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

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Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
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Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400
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Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
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Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
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Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
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Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670
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Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
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Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Gly Ala Pro Gly Thr Pro
        755                 760                 765

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser
    770                 775                 780

Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
785                 790                 795                 800

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
                805                 810                 815

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr
            820                 825                 830

Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
        835                 840                 845

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
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Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly
865                 870                 875                 880

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
                885                 890                 895

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940
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Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
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Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
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Arg Ala Gln Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
            980                 985                 990

Ser Pro Gly Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
        995                 1000                1005

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1010                1015                1020

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
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Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
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Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
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Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1070                1075                1080

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1085                1090                1095

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
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Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
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Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
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Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1145                1150                1155

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
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Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
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Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
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Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
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Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
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Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1235                 1240                 1245

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1250                 1255                 1260

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1265                 1270                 1275

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1280                 1285                 1290

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
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Arg Gly  Ala Pro Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
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Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser
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Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
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Ser Gly  Ser Glu Thr Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
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Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
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Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Pro
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Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Ser Pro  Ala Gly Ser
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Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
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Thr Glu  Glu Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
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Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Ala Ser
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Ser Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
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Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
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Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1490                1495                1500

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1505                1510                1515

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1520                1525                1530

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1535                1540                1545

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1550                1555                1560

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1565                1570                1575

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
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Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1595                1600                1605

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
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Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
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Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
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Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
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Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
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Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
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Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
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His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
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Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1745                1750                1755

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
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Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
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Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
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Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
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Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
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Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
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Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
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Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
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Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    1880                1885                1890

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
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Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
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Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
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Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
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Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    1955                1960                1965
```

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Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
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Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1985                 1990                 1995

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2000                 2005                 2010

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser
    2015                 2020                 2025

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Thr Glu
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Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
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Glu Ser  Gly Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2075                 2080                 2085

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Ser  Glu Pro Ala
    2090                 2095                 2100

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Pro Ala Gly  Ser Pro Thr
    2105                 2110                 2115

Ser Thr  Glu Glu Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
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Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
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Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
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Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu
    2165                 2170                 2175

Gly Ser  Ala Pro Ala Ser Ser  Glu Gln Lys Leu Ile  Ser Glu Glu
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Asp Leu  Ser Pro Ala Thr Gly  His His His His His  His
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
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Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
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Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
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Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
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His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400
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Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            420                 425                 430

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
        435                 440                 445

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    450                 455                 460

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
465                 470                 475                 480

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
                485                 490                 495

Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            500                 505                 510

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
        515                 520                 525

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
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Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670
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Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
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Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
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Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
    930                 935                 940
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Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
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Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
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Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly
            980                 985                 990

Thr Ser Ser Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
        995                 1000                1005

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
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Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser
    1025                1030                1035

Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
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Ser Ser  Thr Gly Ser Pro Ala  Ser Ser Ser Asp Gln  Glu Glu Ile
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Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
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Gln Gly  Ala Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
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Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
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Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1160                1165                1170

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
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Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
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Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
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Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
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Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
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Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
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Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
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Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
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Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
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Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
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Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
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Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
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Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
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Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1370                 1375                 1380

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1385                 1390                 1395

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1400                 1405                 1410

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1415                 1420                 1425

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
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Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
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Arg Gly  Ala Pro Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
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Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser
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Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
    1490                 1495                 1500

Ser Gly  Ser Glu Thr Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1505                 1510                 1515

Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1520                 1525                 1530

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Pro
    1535                 1540                 1545

Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Ser Pro  Ala Gly Ser
    1550                 1555                 1560

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1565                 1570                 1575

Thr Glu  Glu Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1580                 1585                 1590

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Ala Ser
    1595                 1600                 1605

Ser Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1610                 1615                 1620

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1625                 1630                 1635

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1640                 1645                 1650

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1655                 1660                 1665

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1670                 1675                 1680

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1685                 1690                 1695

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1700                 1705                 1710
```

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His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
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Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1730                1735                1740

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1745                1750                1755

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1760                1765                1770

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1775                1780                1785

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1790                1795                1800

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1805                1810                1815

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1820                1825                1830

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1835                1840                1845

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1850                1855                1860

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1865                1870                1875

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1880                1885                1890

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1895                1900                1905

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1910                1915                1920

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1925                1930                1935

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1940                1945                1950

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1955                1960                1965
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Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1970                1975                1980

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1985                1990                1995

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    2000                2005                2010

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    2015                2020                2025

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    2030                2035                2040

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2045                2050                2055

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    2060                2065                2070

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    2075                2080                2085

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2090                2095                2100

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    2105                2110                2115

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    2120                2125                2130

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    2135                2140                2145

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2150                2155                2160

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser
    2165                2170                2175

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Thr Glu
    2180                2185                2190

Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2195                2200                2205

Glu Ser  Gly Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2210                2215                2220
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Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
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Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala
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Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
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Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
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Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
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Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
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Gly Ser Ala Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
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Arg Val Pro Lys Ser Phe Pro Phe Gly Ala Pro Gly Ser Ser Pro Ser
35 40 45

Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
50 55 60

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65 70 75 80

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro
85 90 95

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
100 105 110

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
115 120 125

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser
130 135 140

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145 150 155 160

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
165 170 175

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asn Thr
180 185 190

```
Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460
```

```
His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu
545                 550                 555                 560

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
                565                 570                 575

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            580                 585                 590

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
        595                 600                 605

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    610                 615                 620

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
625                 630                 635                 640

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
                645                 650                 655

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
            660                 665                 670

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
        675                 680                 685

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp
    690                 695                 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735
```

```
Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005
```

```
Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
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Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Gly Ala Pro Ser Pro  Ala Gly Ser
    1040                1045                1050

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1055                1060                1065

Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro
    1070                1075                1080

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1085                1090                1095

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Thr Glu Pro
    1100                1105                1110

Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1115                1120                1125

Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1130                1135                1140

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1145                1150                1155

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1160                1165                1170

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1175                1180                1185

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1190                1195                1200

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1205                1210                1215

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1220                1225                1230

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1235                1240                1245

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1250                1255                1260
```

```
Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1265                1270                1275

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1280                1285                1290

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1295                1300                1305

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1310                1315                1320

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1325                1330                1335

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1340                1345                1350

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1355                1360                1365

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1370                1375                1380

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1385                1390                1395

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1400                1405                1410

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1415                1420                1425

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1430                1435                1440

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1445                1450                1455

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515
```

```
Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                1555                1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                1615                1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770
```

```
Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                1960                1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                1975                1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                1990                1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                2005                2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                2020                2025
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Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
2030 2035 2040

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2045 2050 2055

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
2060 2065 2070

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
2075 2080 2085

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
2090 2095 2100

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
2105 2110 2115

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
2120 2125 2130

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
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Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
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Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Ala Thr Gly His His His His His His
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Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
35 40 45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
50 55 60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65 70 75 80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
85 90 95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
100 105 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
115 120 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
130 135 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145 150 155 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

```
Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
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Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460
```

```
His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu
545                 550                 555                 560

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
                565                 570                 575

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            580                 585                 590

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
        595                 600                 605

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    610                 615                 620

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
625                 630                 635                 640

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
                645                 650                 655

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
            660                 665                 670

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
        675                 680                 685

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp
    690                 695                 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735
```

```
Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005
```

```
Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
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Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Gly Ala Pro Ser Pro  Ala Gly Ser
    1040                1045                1050

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1055                1060                1065

Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro
    1070                1075                1080

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1085                1090                1095

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Thr Glu Pro
    1100                1105                1110

Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1115                1120                1125

Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1130                1135                1140

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1145                1150                1155

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1160                1165                1170

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1175                1180                1185

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1190                1195                1200

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1205                1210                1215

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1220                1225                1230

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1235                1240                1245

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1250                1255                1260
```

```
Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
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Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1280                1285                1290

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1295                1300                1305

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1310                1315                1320

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1325                1330                1335

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1340                1345                1350

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1355                1360                1365

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1370                1375                1380

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1385                1390                1395

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1400                1405                1410

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1415                1420                1425

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1430                1435                1440

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1445                1450                1455

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515
```

```
Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                1555                1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                1615                1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770
```

```
Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                1960                1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                1975                1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                1990                1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                2005                2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                2020                2025
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Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
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Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
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Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
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Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
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Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
2090 2095 2100

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
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Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
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Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
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Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Ala Thr Gly His His His His His His
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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
35 40 45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
50 55 60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65 70 75 80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
85 90 95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
100 105 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
115 120 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
130 135 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145 150 155 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

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Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
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Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460
```

```
His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu
545                 550                 555                 560

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
                565                 570                 575

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            580                 585                 590

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
        595                 600                 605

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    610                 615                 620

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
625                 630                 635                 640

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
                645                 650                 655

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
            660                 665                 670

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
        675                 680                 685

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp
    690                 695                 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735
```

```
Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005
```

```
Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
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Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Gly Ala Pro Ser Pro  Ala Gly Ser
    1040                1045                1050

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1055                1060                1065

Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro
    1070                1075                1080

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1085                1090                1095

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Thr Glu Pro
    1100                1105                1110

Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1115                1120                1125

Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1130                1135                1140

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1145                1150                1155

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1160                1165                1170

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1175                1180                1185

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1190                1195                1200

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1205                1210                1215

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1220                1225                1230

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1235                1240                1245

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1250                1255                1260
```

```
Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1265                1270                1275

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1280                1285                1290

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1295                1300                1305

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1310                1315                1320

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1325                1330                1335

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1340                1345                1350

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1355                1360                1365

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1370                1375                1380

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1385                1390                1395

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1400                1405                1410

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1415                1420                1425

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1430                1435                1440

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1445                1450                1455

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515
```

```
Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                1555                1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                1615                1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770
```

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Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                1960                1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                1975                1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                1990                1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                2005                2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                2020                2025
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Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
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Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
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Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
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Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
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Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
2090 2095 2100

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
2105 2110 2115

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
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Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
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Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
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Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Ala Thr Gly His His His His His His
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35 40 45

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
50 55 60

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
65 70 75 80

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
85 90 95

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
100 105 110

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
115 120 125

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
130 135 140

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
145 150 155 160

Ser Thr Glu Glu Gly Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu

195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
260 265 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
275 280 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
290 295 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305 310 315 320

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325 330 335

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Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
355 360 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
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Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385 390 395 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
405 410 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
420 425 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
435 440 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
450 455 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro

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Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
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Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu
545                 550                 555                 560

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
                565                 570                 575

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            580                 585                 590

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
        595                 600                 605

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    610                 615                 620

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
625                 630                 635                 640

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
                645                 650                 655

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
            660                 665                 670

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
        675                 680                 685

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp
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Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
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Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
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Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
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740 745 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
755 760 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
770 775 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785 790 795 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
805 810 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
820 825 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
835 840 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
850 855 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865 870 875 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
885 890 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
900 905 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
915 920 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
930 935 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945 950 955 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
965 970 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
980 985 990

His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val
995 1000 1005

Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr

1010 1015 1020

Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu
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Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly Ser
1040 1045 1050

Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
1055 1060 1065

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
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Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro
1100 1105 1110

Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
1115 1120 1125

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Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
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Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
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Ser Ala Pro Gly Ala Ser Ser Pro Pro Val Leu Lys Arg His Gln
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Ala Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
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Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
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Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln
1235 1240 1245

Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp
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Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala

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Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu
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Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
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1325 1330 1335

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Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
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1385 1390 1395

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
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Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1415 1420 1425

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
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Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1445 1450 1455

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1460 1465 1470

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1475 1480 1485

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1490 1495 1500

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
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Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn

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Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1535 1540 1545

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
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Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1565 1570 1575

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
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1595 1600 1605

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
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Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1640 1645 1650

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1655 1660 1665

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
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Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1685 1690 1695

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
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Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
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Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
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Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val

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Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
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Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1805 1810 1815

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Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1835 1840 1845

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
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Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1865 1870 1875

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Pro Gly Thr
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Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1940 1945 1950

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1955 1960 1965

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1970 1975 1980

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Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
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Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Gly Ser Ser
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Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
        35                  40                  45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
    50                  55                  60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65                  70                  75                  80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
                85                  90                  95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
            100                 105                 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
        115                 120                 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
    130                 135                 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145                 150                 155                 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205
```

```
Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480
```

```
Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Gly Ala Ser Pro
545                 550                 555                 560

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
                565                 570                 575

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            580                 585                 590

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
        595                 600                 605

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
    610                 615                 620

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
625                 630                 635                 640

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
                645                 650                 655

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
            660                 665                 670

Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
        675                 680                 685

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asp
    690                 695                 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750
```

```
Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020
```

```
Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Gly Ala Pro Ser Pro  Ala Gly Ser
    1040                1045                1050

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1055                1060                1065

Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro
    1070                1075                1080

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1085                1090                1095

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Thr Glu Pro
    1100                1105                1110

Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1115                1120                1125

Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1130                1135                1140

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1145                1150                1155

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1160                1165                1170

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1175                1180                1185

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1190                1195                1200

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1205                1210                1215

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1220                1225                1230

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1235                1240                1245

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1250                1255                1260

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1265                1270                1275
```

```
Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1280                1285                1290

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1295                1300                1305

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1310                1315                1320

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1325                1330                1335

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1340                1345                1350

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1355                1360                1365

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1370                1375                1380

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1385                1390                1395

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1400                1405                1410

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1415                1420                1425

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1430                1435                1440

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1445                1450                1455

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530
```

```
Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                1555                1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                1615                1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785
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Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
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Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
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Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
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Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                1960                1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                1975                1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                1990                1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                2005                2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                2020                2025

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    2030                2035                2040
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Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
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Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
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Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
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Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
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Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
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Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
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Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
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Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
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Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Ala Thr Gly His His His His His His
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Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
35 40 45

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
50 55 60

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
65 70 75 80

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
85 90 95

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
100 105 110

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
115 120 125

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
130 135 140

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
145 150 155 160

Ser Thr Glu Glu Gly Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

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Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
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Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480
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```
Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Gly Ala Ser Pro
545                 550                 555                 560

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
                565                 570                 575

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            580                 585                 590

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
        595                 600                 605

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
    610                 615                 620

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
625                 630                 635                 640

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
                645                 650                 655

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
            660                 665                 670

Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
        675                 680                 685

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asp
    690                 695                 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750
```

```
Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020
```

```
Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Gly Ala Pro Ser Pro  Ala Gly Ser
    1040                1045                1050

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1055                1060                1065

Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro
    1070                1075                1080

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1085                1090                1095

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Thr Glu Pro
    1100                1105                1110

Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1115                1120                1125

Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1130                1135                1140

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1145                1150                1155

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1160                1165                1170

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1175                1180                1185

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1190                1195                1200

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1205                1210                1215

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1220                1225                1230

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1235                1240                1245

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1250                1255                1260

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1265                1270                1275
```

```
Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1280                 1285                 1290

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1295                 1300                 1305

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1310                 1315                 1320

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1325                 1330                 1335

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1340                 1345                 1350

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1355                 1360                 1365

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1370                 1375                 1380

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1385                 1390                 1395

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1400                 1405                 1410

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1415                 1420                 1425

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1430                 1435                 1440

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1445                 1450                 1455

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                 1465                 1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                 1480                 1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                 1495                 1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                 1510                 1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                 1525                 1530
```

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Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                1555                1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                1615                1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
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Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785
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Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
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Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
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Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
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Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
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Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                1990                1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                2005                2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                2020                2025

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
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Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
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Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
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Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
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Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
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Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
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Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
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Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
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Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
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Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Arg Val Pro Lys Ser Phe Pro Phe Gly Ala Pro Gly Ser Ser Pro Ser
35 40 45

Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
50 55 60

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65 70 75 80

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro
85 90 95

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
100 105 110

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
115 120 125

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser
130 135 140

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145 150 155 160

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
165 170 175

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

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Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480
```

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Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Gly Ala Ser Pro
545                 550                 555                 560

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
                565                 570                 575

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            580                 585                 590

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
        595                 600                 605

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
    610                 615                 620

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
625                 630                 635                 640

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
                645                 650                 655

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
            660                 665                 670

Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
        675                 680                 685

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asp
    690                 695                 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750
```

```
Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020
```

```
Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
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Pro Arg  Ser Phe Ser Gln Asn  Gly Ala Pro Ser Pro  Ala Gly Ser
    1040                1045                1050

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1055                1060                1065

Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro
    1070                1075                1080

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1085                1090                1095

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Thr Glu Pro
    1100                1105                1110

Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1115                1120                1125

Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1130                1135                1140

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1145                1150                1155

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1160                1165                1170

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1175                1180                1185

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1190                1195                1200

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1205                1210                1215

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1220                1225                1230

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1235                1240                1245

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1250                1255                1260

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1265                1270                1275
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Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1280                1285                1290

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1295                1300                1305

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1310                1315                1320

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1325                1330                1335

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1340                1345                1350

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1355                1360                1365

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1370                1375                1380

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1385                1390                1395

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1400                1405                1410

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1415                1420                1425

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1430                1435                1440

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1445                1450                1455

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530
```

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Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                1555                1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                1615                1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
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Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785
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Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
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Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
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Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
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Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                1960                1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                1975                1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                1990                1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                2005                2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                2020                2025

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    2030                2035                2040
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Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
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Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
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Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
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Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
35 40 45

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
50 55 60

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
65 70 75 80

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
85 90 95

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
100 105 110

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
115 120 125

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
130 135 140

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
145 150 155 160

Ser Thr Glu Glu Gly Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro

210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
260 265 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
275 280 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
290 295 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305 310 315 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
325 330 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
340 345 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
355 360 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
370 375 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385 390 395 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
405 410 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
420 425 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
435 440 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
450 455 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465 470 475 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr

485 490 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
500 505 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
515 520 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
530 535 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gly
545 550 555 560

Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro
565 570 575

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
580 585 590

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
595 600 605

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
610 615 620

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
625 630 635 640

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
645 650 655

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
660 665 670

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
675 680 685

Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
690 695 700

Pro Gly Ala Ser Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705 710 715 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
725 730 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
740 745 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln

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        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
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Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
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Pro Arg  Ser Phe Ser Gln Asn  Gly Ala Pro Ser Pro  Ala Gly Ser
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Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
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Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro
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Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1085                1090                1095

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Thr Glu Pro
    1100                1105                1110

Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
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Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
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Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1145                1150                1155

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1160                1165                1170

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1175                1180                1185

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1190                1195                1200

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1205                1210                1215

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1220                1225                1230

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1235                1240                1245

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1250                1255                1260

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1265                1270                1275

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
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Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
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Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1310 1315 1320

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1325 1330 1335

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1340 1345 1350

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1355 1360 1365

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1370 1375 1380

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1385 1390 1395

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1400 1405 1410

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1415 1420 1425

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1430 1435 1440

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1445 1450 1455

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1460 1465 1470

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1475 1480 1485

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1490 1495 1500

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1505 1510 1515

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1520 1525 1530

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys

1535 1540 1545

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1550 1555 1560

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1565 1570 1575

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1580 1585 1590

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1595 1600 1605

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
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Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
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Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1640 1645 1650

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1655 1660 1665

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1670 1675 1680

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1685 1690 1695

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1700 1705 1710

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1715 1720 1725

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1730 1735 1740

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
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Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1760 1765 1770

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
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Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly

1790 1795 1800

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1805 1810 1815

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1820 1825 1830

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1835 1840 1845

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1850 1855 1860

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1865 1870 1875

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Pro Gly Thr
1880 1885 1890

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1895 1900 1905

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
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Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
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Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1940 1945 1950

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
1955 1960 1965

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
1970 1975 1980

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1985 1990 1995

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2000 2005 2010

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
2015 2020 2025

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
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2075 2080 2085

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Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Ala Thr Gly His His His His His His
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tcggaaacgc caggcacaag tgagtctgca actcccgagt ccggacctgg ctccgagcct 180

gccactagcg gctccgagac tccgggaact tccgagagcg ctacaccaga aagcggaccc 240

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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
35 40 45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
50 55 60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65 70 75 80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
85 90 95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
100 105 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
115 120 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
130 135 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145 150 155 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

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Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495
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Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gly
545                 550                 555                 560

Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro
                565                 570                 575

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
            580                 585                 590

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
        595                 600                 605

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
    610                 615                 620

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
625                 630                 635                 640

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
                645                 650                 655

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
            660                 665                 670

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
        675                 680                 685

Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    690                 695                 700

Pro Gly Ala Ser Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765
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Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020

Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
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Pro Arg  Ser Phe Ser Gln Asn  Gly Ala Pro Ser Pro  Ala Gly Ser
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Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
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Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro
    1070                1075                1080

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1085                1090                1095

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Thr Glu Pro
    1100                1105                1110

Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1115                1120                1125

Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1130                1135                1140

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1145                1150                1155

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1160                1165                1170

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1175                1180                1185

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1190                1195                1200

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1205                1210                1215

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1220                1225                1230

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1235                1240                1245

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1250                1255                1260

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1265                1270                1275

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1280                1285                1290
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Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
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Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
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Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1325                1330                1335

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
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Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
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Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
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Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
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Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
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Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
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Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1430                1435                1440

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1445                1450                1455

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
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Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
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Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
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Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
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Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
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Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
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Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
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Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
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Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
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Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
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Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
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Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
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Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
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Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
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Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
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Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
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Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
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Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
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Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
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Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
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Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
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Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
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Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
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Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
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Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
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Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
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Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
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<223> Description of Artificial Sequence: Synthetic polynucleotide

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cccggaacgt catcgaccgg atcgccaggg agctcgacgc catcaggcgc aacaggatca 240

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agcaccggat cacccggatc gagcccatcc gcctcaaccg gaaccggacc cggtacacca 360

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cccggttcgt caacgccttc cggagcaact ggtagccccg gagcatcgcc aggaacttcg 480

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accgagccct ccgaaggatc agccccaggc accagcacgg agccctctga gggaagcgca 2040

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<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Gly Ser Ser
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Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
35 40 45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
50 55 60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65 70 75 80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
85 90 95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
100 105 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
115 120 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
130 135 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145 150 155 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

```
Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495
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Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gly
545                 550                 555                 560

Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro
                565                 570                 575

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
            580                 585                 590

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
        595                 600                 605

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
    610                 615                 620

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
625                 630                 635                 640

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
                645                 650                 655

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
            660                 665                 670

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
        675                 680                 685

Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    690                 695                 700

Pro Gly Ala Ser Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765
```

```
Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020

Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035
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Pro Arg  Ser Phe Ser Gln Asn  Gly Ala Pro Ser Pro  Ala Gly Ser
    1040                1045                1050

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1055                1060                1065

Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro
    1070                1075                1080

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1085                1090                1095

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Thr Glu Pro
    1100                1105                1110

Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1115                1120                1125

Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1130                1135                1140

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1145                1150                1155

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1160                1165                1170

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1175                1180                1185

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1190                1195                1200

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1205                1210                1215

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1220                1225                1230

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1235                1240                1245

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1250                1255                1260

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1265                1270                1275

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1280                1285                1290
```

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Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1295                1300                1305

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1310                1315                1320

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1325                1330                1335

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1340                1345                1350

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1355                1360                1365

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1370                1375                1380

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1385                1390                1395

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1400                1405                1410

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1415                1420                1425

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1430                1435                1440

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1445                1450                1455

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545
```

```
Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                1555                1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                1615                1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800
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Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                 1810                 1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                 1825                 1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                 1840                 1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                 1855                 1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                 1870                 1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                 1885                 1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                 1900                 1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                 1915                 1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                 1930                 1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                 1945                 1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                 1960                 1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                 1975                 1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                 1990                 1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                 2005                 2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                 2020                 2025

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    2030                 2035                 2040

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2045                 2050                 2055
```

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
2060 2065 2070

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
2075 2080 2085

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
2090 2095 2100

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
2105 2110 2115

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
2120 2125 2130

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
2135 2140 2145

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2150 2155 2160

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2165 2170 2175

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
2180 2185 2190

Pro Ala Thr Gly His His His His His His
2195 2200

<210> 1222
<211> 6612
<212> DNA
<213> Artificial Sequence

<220>
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ccaggttcgt caccctcagc atcaacgggg acgggaccgg gggcgtcacc aggaacgtcc 180

tccaccggct cgccgggtgc atcacccgga acgtcatcga ccggatcgcc agggagctcg 240

acgccatcag gcgcaacagg atcacctggc tcaagcccta gcgcgtcaac cggcacgggt 300

ccgggtgcct cccctggcac gtccagcacc ggatcacccg gatcgagccc atccgcctca 360

accggaaccg gacccggtac accagggtcg ggaacagcct cctcgtcacc aggctcctca 420

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cccggagcat cgccaggaac ttcgagcacg gggtcgcccg cctcgagcga cgcaagattt 540

cctcctagag tgccaaaatc ttttccattc aacacctcag tcgtgtacaa aaagactctg 600

tttgtagaat tcacggatca ccttttcaac atcgctaagc caaggccacc ctggatgggt 660

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atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag 780

ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct 840

ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

ggcctcattg gagccctact agtatgtaga gaagggagtc tggccaagga aaagacacag 1020

accttgcaca aatttatact acttttgct gtatttgatg aagggaaaag ttggcactca 1080

gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg 1200

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ataactttcc ttactgctca aacactcttg atggaccttg gacagtttct actgttttgt 1380

catatctctt cccaccaaca tgatggcatg gaagcttatg tcaaagtaga cagctgtcca 1440

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actgattctg aaatggatgt ggtcaggttt gatgatgaca actctccttc ctttatccaa 1560

attcgctcag ttgccaagaa gcatcctaaa acttgggtac attacattgc tgctgaagag 1620

gaggactggg actatgctcc cttagtcctc gcccccgatg acagaagtta taaaagtggc 1680

gcgccaacca gtacggagcc gtccgagggg agcgcaccag gaagcccggc tgggagcccg 1740

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<210> 1223
<211> 2203
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Gly Ala Pro Gly Ser Ser Pro Ser
35 40 45

Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
50 55 60

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65 70 75 80

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro
85 90 95

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
100 105 110

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
115 120 125

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser
130 135 140

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145 150 155 160

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
165 170 175

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

```
Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495
```

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Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gly
545                 550                 555                 560

Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro
                565                 570                 575

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
            580                 585                 590

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
        595                 600                 605

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
    610                 615                 620

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
625                 630                 635                 640

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
                645                 650                 655

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
            660                 665                 670

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
        675                 680                 685

Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    690                 695                 700

Pro Gly Ala Ser Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765
```

```
Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020

Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035
```

```
Pro Arg  Ser Phe Ser Gln Asn  Gly Ala Pro Ser Pro  Ala Gly Ser
    1040                1045                1050

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1055                1060                1065

Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro
    1070                1075                1080

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1085                1090                1095

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Thr Glu Pro
    1100                1105                1110

Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1115                1120                1125

Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1130                1135                1140

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1145                1150                1155

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1160                1165                1170

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1175                1180                1185

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1190                1195                1200

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1205                1210                1215

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1220                1225                1230

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1235                1240                1245

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1250                1255                1260

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1265                1270                1275

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1280                1285                1290
```

```
Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1295                1300                1305

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1310                1315                1320

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1325                1330                1335

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1340                1345                1350

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1355                1360                1365

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1370                1375                1380

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1385                1390                1395

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1400                1405                1410

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1415                1420                1425

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1430                1435                1440

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1445                1450                1455

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545
```

```
Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                1555                1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                1615                1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800
```

```
Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                 1810                 1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                 1825                 1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                 1840                 1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                 1855                 1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                 1870                 1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                 1885                 1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                 1900                 1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                 1915                 1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                 1930                 1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                 1945                 1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                 1960                 1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                 1975                 1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                 1990                 1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                 2005                 2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                 2020                 2025

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    2030                 2035                 2040

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2045                 2050                 2055
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Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
2060 2065 2070

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
2075 2080 2085

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
2090 2095 2100

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
2105 2110 2115

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
2120 2125 2130

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
2135 2140 2145

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2150 2155 2160

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2165 2170 2175

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
2180 2185 2190

Pro Ala Thr Gly His His His His His His
2195 2200

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<211> 6612
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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ggcgcgccag gctccagccc ctccgcgagc acgggaaccg gaccaggttc gtcaccctca 180

gcatcaacgg ggacgggacc gggggcgtca ccaggaacgt cctccaccgg ctcgccgggt 240

gcatcacccg gaacgtcatc gaccggatcg ccagggagct cgacgccatc aggcgcaaca 300

ggatcacctg gctcaagccc tagcgcgtca accggcacgg gtccgggtgc ctcccctggc 360

acgtccagca ccggatcacc cggatcgagc ccatccgcct caaccggaac cggacccggt 420

acaccagggt cgggaacagc ctcctcgtca ccaggctcct caacccctc gggagccacg 480

ggttcgcccg gttcgtcaac gccttccgga gcaactggta gccccggagc atcgccagga 540

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tttgtagaat tcacggatca ccttttcaac atcgctaagc caaggccacc ctggatgggt 660

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atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag 780

ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct 840

ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

ggcctcattg gagccctact agtatgtaga gaagggagtc tggccaagga aaagacacag 1020

accttgcaca aatttatact acttttgct gtatttgatg aagggaaaag ttggcactca 1080

gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg 1200

aaatcagtct attggcatgt gattggaatg ggcaccactc ctgaagtgca ctcaatattc 1260

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gccacccccg agagtggccc aggcactagt accgagccct ctgaaggcag tgcgccagcc 6540

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catcaccatt ga 6612

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<211> 2203
<212> PRT
<213> Artificial Sequence

<220>
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Gly Ala Pro Gly Ser Ser Pro Ser
35 40 45

Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
50 55 60

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65 70 75 80

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro
85 90 95

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
100 105 110

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
115 120 125

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser
130 135 140

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145 150 155 160

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
165 170 175

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn

225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
260 265 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
275 280 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
290 295 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305 310 315 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
325 330 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
340 345 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
355 360 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
370 375 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385 390 395 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
405 410 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
420 425 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
435 440 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
450 455 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465 470 475 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
485 490 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp

500 505 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
515 520 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
530 535 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gly
545 550 555 560

Ala Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser
565 570 575

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala
580 585 590

Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
595 600 605

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser
610 615 620

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr
625 630 635 640

Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
645 650 655

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr
660 665 670

Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr
675 680 685

Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
690 695 700

Ser Pro Ala Ser Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705 710 715 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
725 730 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
740 745 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
755 760 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg

770 775 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785 790 795 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
805 810 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
820 825 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
835 840 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
850 855 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865 870 875 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
885 890 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
900 905 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
915 920 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
930 935 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945 950 955 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
965 970 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
980 985 990

His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val
995 1000 1005

Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr
1010 1015 1020

Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu
1025 1030 1035

Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly Ser

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Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1055                1060                1065

Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro
    1070                1075                1080

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1085                1090                1095

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Thr Glu Pro
    1100                1105                1110

Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1115                1120                1125

Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1130                1135                1140

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1145                1150                1155

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1160                1165                1170

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1175                1180                1185

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1190                1195                1200

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1205                1210                1215

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1220                1225                1230

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1235                1240                1245

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1250                1255                1260

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1265                1270                1275

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1280                1285                1290

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
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1295 1300 1305

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1310 1315 1320

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1325 1330 1335

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1340 1345 1350

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1355 1360 1365

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1370 1375 1380

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1385 1390 1395

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1400 1405 1410

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1415 1420 1425

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1430 1435 1440

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1445 1450 1455

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1460 1465 1470

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1475 1480 1485

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1490 1495 1500

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1505 1510 1515

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1520 1525 1530

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1535 1540 1545

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His

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Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
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Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                1615                1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
```

1805 1810 1815

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1820 1825 1830

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1835 1840 1845

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1850 1855 1860

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1865 1870 1875

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Pro Gly Thr
1880 1885 1890

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1895 1900 1905

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
1910 1915 1920

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1925 1930 1935

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1940 1945 1950

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
1955 1960 1965

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
1970 1975 1980

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1985 1990 1995

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2000 2005 2010

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
2015 2020 2025

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
2030 2035 2040

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2045 2050 2055

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr

2060 2065 2070

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
2075 2080 2085

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
2090 2095 2100

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
2105 2110 2115

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
2120 2125 2130

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
2135 2140 2145

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2150 2155 2160

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2165 2170 2175

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
2180 2185 2190

Pro Ala Thr Gly His His His His His His
2195 2200

<210> 1226
<211> 6612
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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ggcgcgccag gctccagccc ctccgcgagc acgggaaccg gaccaggttc gtcaccctca 180

gcatcaacgg ggacgggacc gggggcgtca ccaggaacgt cctccaccgg ctcgccgggt 240

gcatcacccg gaacgtcatc gaccggatcg ccagggagct cgacgccatc aggcgcaaca 300

ggatcacctg gctcaagccc tagcgcgtca accggcacgg gtccgggtgc ctcccctggc 360

acgtccagca ccggatcacc cggatcgagc ccatccgcct caaccggaac cggacccggt 420

acaccagggt cgggaacagc ctcctcgtca ccaggctcct caacccctc gggagccacg 480

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<210> 1227
<211> 2203
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1227

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Met Gln Gly Ala Pro Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
            20                  25                  30

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
        35                  40                  45

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
    50                  55                  60

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
65                  70                  75                  80

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
                85                  90                  95

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
            100                 105                 110

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
        115                 120                 125

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
    130                 135                 140

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
145                 150                 155                 160

Ser Thr Glu Glu Gly Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240
```

```
Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510
```

```
Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gly
545                 550                 555                 560

Ala Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser
                565                 570                 575

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala
            580                 585                 590

Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
        595                 600                 605

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser
    610                 615                 620

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr
625                 630                 635                 640

Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
                645                 650                 655

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr
            660                 665                 670

Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr
        675                 680                 685

Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
    690                 695                 700

Ser Pro Ala Ser Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780
```

```
Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020

Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Gly Ala Pro Ser Pro  Ala Gly Ser
    1040                1045                1050
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```
Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1055                 1060                 1065

Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro
    1070                 1075                 1080

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1085                 1090                 1095

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Thr Glu Pro
    1100                 1105                 1110

Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1115                 1120                 1125

Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1130                 1135                 1140

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1145                 1150                 1155

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1160                 1165                 1170

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1175                 1180                 1185

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1190                 1195                 1200

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1205                 1210                 1215

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1220                 1225                 1230

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1235                 1240                 1245

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1250                 1255                 1260

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1265                 1270                 1275

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1280                 1285                 1290

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1295                 1300                 1305
```

```
Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1310                1315                1320

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1325                1330                1335

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1340                1345                1350

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1355                1360                1365

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1370                1375                1380

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1385                1390                1395

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1400                1405                1410

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1415                1420                1425

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1430                1435                1440

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1445                1450                1455

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                1555                1560
```

```
Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                1615                1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815
```

```
Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                 1825                 1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                 1840                 1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                 1855                 1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                 1870                 1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                 1885                 1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                 1900                 1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                 1915                 1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                 1930                 1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                 1945                 1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                 1960                 1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                 1975                 1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                 1990                 1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                 2005                 2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                 2020                 2025

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    2030                 2035                 2040

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2045                 2050                 2055

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    2060                 2065                 2070
```

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
2075 2080 2085

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
2090 2095 2100

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
2105 2110 2115

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
2120 2125 2130

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
2135 2140 2145

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2150 2155 2160

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2165 2170 2175

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
2180 2185 2190

Pro Ala Thr Gly His His His His His His
2195 2200

<210> 1228
<211> 6612
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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tcggaaacgc caggcacaag tgagtctgca actcccgagt ccggacctgg ctccgagcct 180

gccactagcg gctccgagac tccgggaact tccgagagcg ctacaccaga aagcggaccc 240

ggaaccagta ccgaacctag cgagggctct gctccgggca gcccagccgg ctctcctaca 300

tccacggagg agggcacttc cgaatccgcc accccggagt cagggccagg atctgaaccc 360

gctacctcag gcagtgagac gccaggaacg agcgagtccg ctacaccgga gagtgggcca 420

gggagccctg ctggatctcc tacgtccact gaggaagggt caccagcggg ctcgcccacc 480

agcactgaag aaggtgcctc gagcagtgat ctcggtgagc tgcctgtgga cgcaagattt 540

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tttgtagaat tcacggatca ccttttcaac atcgctaagc caaggccacc ctggatgggt 660

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atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag 780

ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct 840

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ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

ggcctcattg gagccctact agtatgtaga gaagggagtc tggccaagga aaagacacag 1020

accttgcaca aatttatact acttttgct gtatttgatg aagggaaaag ttggcactca 1080

gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

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aaatcagtct attggcatgt gattggaatg ggcaccactc ctgaagtgca ctcaatattc 1260

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ataactttcc ttactgctca aacactcttg atggaccttg gacagtttct actgttttgt 1380

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tcagcacccg gctctgagcc ggccacaagt ggcagtgaga cacccggcac ttcagagagt 6480

gccacccccg agagtggccc aggcactagt accgagccct ctgaaggcag tgcgccagcc 6540

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<210> 1229

<211> 2203
<212> PRT
<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1229

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Gly Ser Ser
            20                  25                  30

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
        35                  40                  45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
    50                  55                  60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65                  70                  75                  80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
                85                  90                  95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
            100                 105                 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
        115                 120                 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
    130                 135                 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145                 150                 155                 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240
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```
Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510
```

```
Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gly
545                 550                 555                 560

Ala Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser
                565                 570                 575

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala
            580                 585                 590

Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
        595                 600                 605

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser
    610                 615                 620

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr
625                 630                 635                 640

Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
                645                 650                 655

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr
            660                 665                 670

Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr
        675                 680                 685

Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
    690                 695                 700

Ser Pro Ala Ser Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750

Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780
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```
Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020

Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Gly Ala Pro Ser Pro  Ala Gly Ser
    1040                1045                1050
```

```
Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1055                 1060                 1065

Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro
    1070                 1075                 1080

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1085                 1090                 1095

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Thr Glu Pro
    1100                 1105                 1110

Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1115                 1120                 1125

Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1130                 1135                 1140

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1145                 1150                 1155

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1160                 1165                 1170

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1175                 1180                 1185

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1190                 1195                 1200

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1205                 1210                 1215

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1220                 1225                 1230

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1235                 1240                 1245

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1250                 1255                 1260

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1265                 1270                 1275

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1280                 1285                 1290

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1295                 1300                 1305
```

```
Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1310                1315                1320

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1325                1330                1335

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1340                1345                1350

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1355                1360                1365

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1370                1375                1380

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1385                1390                1395

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1400                1405                1410

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1415                1420                1425

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
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Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1445                1450                1455

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                1555                1560
```

```
Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                1615                1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815
```

```
Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                 1825                 1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                 1840                 1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                 1855                 1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                 1870                 1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                 1885                 1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                 1900                 1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                 1915                 1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                 1930                 1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                 1945                 1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                 1960                 1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                 1975                 1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                 1990                 1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                 2005                 2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                 2020                 2025

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    2030                 2035                 2040

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2045                 2050                 2055

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    2060                 2065                 2070
```

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
2075 2080 2085

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
2090 2095 2100

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
2105 2110 2115

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
2120 2125 2130

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
2135 2140 2145

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2150 2155 2160

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2165 2170 2175

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
2180 2185 2190

Pro Ala Thr Gly His His His His His His
2195 2200

<210> 1230
<211> 6612
<212> DNA
<213> Artificial Sequence

<220>
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ccaggttcgt caccctcagc atcaacgggg acgggaccgg gggcgtcacc aggaacgtcc 180

tccaccggct cgccgggtgc atcacccgga acgtcatcga ccggatcgcc agggagctcg 240

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<210> 1231
<211> 2206

<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1231

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Gly Ala Pro Gly Ser Ser Pro Ser
        35                  40                  45

Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
    50                  55                  60

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65                  70                  75                  80

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro
                85                  90                  95

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
            100                 105                 110

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
        115                 120                 125

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser
    130                 135                 140

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145                 150                 155                 160

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
                165                 170                 175

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240
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```
Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510
```

```
Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780
```

```
Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050
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Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
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Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Gly  Ala Pro Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1130                1135                1140

Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser
    1145                1150                1155

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
    1160                1165                1170

Ser Gly  Ser Glu Thr Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1175                1180                1185

Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1190                1195                1200

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Pro
    1205                1210                1215

Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Ser Pro  Ala Gly Ser
    1220                1225                1230

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
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Thr Glu  Glu Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1250                1255                1260

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Ala Ser
    1265                1270                1275

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1280                1285                1290

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1295                1300                1305
```

```
Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
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Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1325                 1330                 1335

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1340                 1345                 1350

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1355                 1360                 1365

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1370                 1375                 1380

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1385                 1390                 1395

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1400                 1405                 1410

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
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Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1430                 1435                 1440

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1445                 1450                 1455

Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1460                 1465                 1470

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1475                 1480                 1485

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1490                 1495                 1500

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1505                 1510                 1515

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1520                 1525                 1530

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1535                 1540                 1545

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1550                 1555                 1560
```

```
His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
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Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1580                1585                1590

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1595                1600                1605

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1610                1615                1620

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1625                1630                1635

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1640                1645                1650

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1655                1660                1665

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1670                1675                1680

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1685                1690                1695

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1700                1705                1710

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1715                1720                1725

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1730                1735                1740

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1745                1750                1755

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1760                1765                1770

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1775                1780                1785

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1790                1795                1800

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1805                1810                1815
```

```
Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1820                1825                1830

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1835                1840                1845

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    1850                1855                1860

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    1865                1870                1875

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    1880                1885                1890

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1895                1900                1905

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    1910                1915                1920

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1925                1930                1935

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1940                1945                1950

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    1955                1960                1965

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    1970                1975                1980

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1985                1990                1995

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2000                2005                2010

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser
    2015                2020                2025

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Thr Glu
    2030                2035                2040

Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2045                2050                2055

Glu Ser  Gly Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2060                2065                2070
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Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2075 2080 2085

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala
2090 2095 2100

Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
2105 2110 2115

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2120 2125 2130

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
2135 2140 2145

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
2150 2155 2160

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
2165 2170 2175

Gly Ser Ala Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu
2180 2185 2190

Asp Leu Ser Pro Ala Thr Gly His His His His His His
2195 2200 2205

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<211> 6621
<212> DNA
<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic polynucleotide

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gcatcaacgg ggacgggacc gggggcgtca ccaggaacgt cctccaccgg ctcgccgggt 240

gcatcacccg gaacgtcatc gaccggatcg ccagggagct cgacgccatc aggcgcaaca 300

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<211> 2206
<212> PRT

<213> Artificial Sequence

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Gly Ser Ser
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Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
35 40 45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
50 55 60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65 70 75 80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
85 90 95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
100 105 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
115 120 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
130 135 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145 150 155 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp

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                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
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Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
530 535 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545 550 555 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
565 570 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
580 585 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
595 600 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
610 615 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
645 650 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser

785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
835 840 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
885 890 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
900 905 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
915 920 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
930 935 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945 950 955 960

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Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
980 985 990

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
995 1000 1005

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser
1010 1015 1020

Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
1025 1030 1035

Ser Ala Pro Gly Ala Ser Ser Pro Pro Val Leu Lys Arg His Gln
1040 1045 1050

Ala Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile

1055 1060 1065

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
1070 1075 1080

Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln
1085 1090 1095

Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp
1100 1105 1110

Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala
1115 1120 1125

Gln Gly Ala Pro Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
1130 1135 1140

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser
1145 1150 1155

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr
1160 1165 1170

Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
1175 1180 1185

Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
1190 1195 1200

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro
1205 1210 1215

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser
1220 1225 1230

Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser
1235 1240 1245

Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
1250 1255 1260

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser
1265 1270 1275

Ser Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val
1280 1285 1290

Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg
1295 1300 1305

Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg

1310 1315 1320

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala
1325 1330 1335

Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu
1340 1345 1350

Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro
1355 1360 1365

Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala
1370 1375 1380

Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser
1385 1390 1395

Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
1400 1405 1410

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg
1415 1420 1425

Gln Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp
1430 1435 1440

Glu Thr Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys
1445 1450 1455

Arg Ala Pro Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu
1460 1465 1470

Asn Tyr Arg Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu
1475 1480 1485

Pro Gly Leu Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu
1490 1495 1500

Leu Ser Met Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser
1505 1510 1515

Gly His Val Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala
1520 1525 1530

Leu Tyr Asn Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu
1535 1540 1545

Pro Ser Lys Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu
1550 1555 1560

His Leu His Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn

1565 1570 1575

Lys Cys Gln Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp
1580 1585 1590

Phe Gln Ile Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys
1595 1600 1605

Leu Ala Arg Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr
1610 1615 1620

Lys Glu Pro Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met
1625 1630 1635

Ile Ile His Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser
1640 1645 1650

Ser Leu Tyr Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly
1655 1660 1665

Lys Lys Trp Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met
1670 1675 1680

Val Phe Phe Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile
1685 1690 1695

Phe Asn Pro Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr
1700 1705 1710

His Tyr Ser Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys
1715 1720 1725

Asp Leu Asn Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala
1730 1735 1740

Ile Ser Asp Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met
1745 1750 1755

Phe Ala Thr Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly
1760 1765 1770

Arg Ser Asn Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp
1775 1780 1785

Leu Gln Val Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr
1790 1795 1800

Thr Gln Gly Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu
1805 1810 1815

Phe Leu Ile Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe

1820 1825 1830

Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser
1835 1840 1845

Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg
1850 1855 1860

Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu
1865 1870 1875

Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala
1880 1885 1890

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
1895 1900 1905

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
1910 1915 1920

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
1925 1930 1935

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1940 1945 1950

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
1955 1960 1965

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
1970 1975 1980

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
1985 1990 1995

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2000 2005 2010

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser
2015 2020 2025

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu
2030 2035 2040

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
2045 2050 2055

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2060 2065 2070

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser

2075 2080 2085

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala
2090 2095 2100

Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
2105 2110 2115

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2120 2125 2130

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
2135 2140 2145

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
2150 2155 2160

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
2165 2170 2175

Gly Ser Ala Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu
2180 2185 2190

Asp Leu Ser Pro Ala Thr Gly His His His His His His
2195 2200 2205

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<211> 6621
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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tccaccggct cgccgggtgc atcacccgga acgtcatcga ccggatcgcc agggagctcg 240

acgccatcag gcgcaacagg atcacctggc tcaagcccta gcgcgtcaac cggcacgggt 300

ccgggtgcct cccctggcac gtccagcacc ggatcacccg gatcgagccc atccgcctca 360

accggaaccg gacccggtac accagggtcg ggaacagcct cctcgtcacc aggctcctca 420

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ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

ggcctcattg gagccctact agtatgtaga gaagggagtc tggccaagga aaagacacag 1020

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<211> 2206
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1235

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            20                  25                  30

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
        35                  40                  45

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
    50                  55                  60

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
65                  70                  75                  80

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
                85                  90                  95

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
            100                 105                 110

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
        115                 120                 125

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
    130                 135                 140

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
145                 150                 155                 160

Ser Thr Glu Glu Gly Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255
```

```
Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525
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```
Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800
```

```
Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065
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Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
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Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Gly  Ala Pro Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1130                1135                1140

Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser
    1145                1150                1155

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
    1160                1165                1170

Ser Gly  Ser Glu Thr Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1175                1180                1185

Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1190                1195                1200

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Pro
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Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Ser Pro  Ala Gly Ser
    1220                1225                1230

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1235                1240                1245

Thr Glu  Glu Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1250                1255                1260

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Ala Ser
    1265                1270                1275

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1280                1285                1290

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1295                1300                1305

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1310                1315                1320
```

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Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
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Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
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Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1355                1360                1365

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1370                1375                1380

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1385                1390                1395

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1400                1405                1410

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1415                1420                1425

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1430                1435                1440

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1445                1450                1455

Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1460                1465                1470

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1475                1480                1485

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1490                1495                1500

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1505                1510                1515

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1520                1525                1530

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1535                1540                1545

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1550                1555                1560

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1565                1570                1575
```

```
Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1580                 1585                 1590

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1595                 1600                 1605

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1610                 1615                 1620

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1625                 1630                 1635

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1640                 1645                 1650

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1655                 1660                 1665

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1670                 1675                 1680

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1685                 1690                 1695

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1700                 1705                 1710

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1715                 1720                 1725

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1730                 1735                 1740

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1745                 1750                 1755

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1760                 1765                 1770

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1775                 1780                 1785

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1790                 1795                 1800

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1805                 1810                 1815

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1820                 1825                 1830
```

```
Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1835                 1840                 1845

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    1850                 1855                 1860

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    1865                 1870                 1875

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    1880                 1885                 1890

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1895                 1900                 1905

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    1910                 1915                 1920

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1925                 1930                 1935

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1940                 1945                 1950

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    1955                 1960                 1965

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    1970                 1975                 1980

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1985                 1990                 1995

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2000                 2005                 2010

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser
    2015                 2020                 2025

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Thr Glu
    2030                 2035                 2040

Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2045                 2050                 2055

Glu Ser  Gly Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2060                 2065                 2070

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2075                 2080                 2085
```

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala
2090 2095 2100

Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
2105 2110 2115

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2120 2125 2130

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
2135 2140 2145

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
2150 2155 2160

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
2165 2170 2175

Gly Ser Ala Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu
2180 2185 2190

Asp Leu Ser Pro Ala Thr Gly His His His His His His
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<211> 6621
<212> DNA
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gccactagcg gctccgagac tccgggaact tccgagagcg ctacaccaga aagcggaccc 240

ggaaccagta ccgaacctag cgagggctct gctccgggca gcccagccgg ctctcctaca 300

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gctacctcag gcagtgagac gccaggaacg agcgagtccg ctacaccgga gagtgggcca 420

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<212> PRT
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<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1237

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            20                  25                  30

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
        35                  40                  45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
    50                  55                  60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65                  70                  75                  80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
                85                  90                  95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
            100                 105                 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
        115                 120                 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
    130                 135                 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145                 150                 155                 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255
```

```
Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525
```

```
Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800
```

```
Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
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Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
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Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
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Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065
```

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Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
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Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
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Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Gly  Ala Pro Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1130                1135                1140

Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser
    1145                1150                1155

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
    1160                1165                1170

Ser Gly  Ser Glu Thr Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
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Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
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Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Pro
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Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Ser Pro  Ala Gly Ser
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Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1235                1240                1245

Thr Glu  Glu Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
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Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Ala Ser
    1265                1270                1275

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1280                1285                1290

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1295                1300                1305

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
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Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
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Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
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Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1355                1360                1365

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1370                1375                1380

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1385                1390                1395

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1400                1405                1410

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1415                1420                1425

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1430                1435                1440

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1445                1450                1455

Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1460                1465                1470

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1475                1480                1485

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1490                1495                1500

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1505                1510                1515

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1520                1525                1530

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1535                1540                1545

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1550                1555                1560

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1565                1570                1575
```

```
Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1580                 1585                 1590

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1595                 1600                 1605

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1610                 1615                 1620

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1625                 1630                 1635

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1640                 1645                 1650

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1655                 1660                 1665

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1670                 1675                 1680

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1685                 1690                 1695

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1700                 1705                 1710

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1715                 1720                 1725

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1730                 1735                 1740

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1745                 1750                 1755

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1760                 1765                 1770

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1775                 1780                 1785

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1790                 1795                 1800

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1805                 1810                 1815

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1820                 1825                 1830
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Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
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Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    1850                1855                1860

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    1865                1870                1875

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    1880                1885                1890

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1895                1900                1905

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    1910                1915                1920

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1925                1930                1935

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1940                1945                1950

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    1955                1960                1965

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    1970                1975                1980

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1985                1990                1995

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2000                2005                2010

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser
    2015                2020                2025

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Thr Glu
    2030                2035                2040

Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2045                2050                2055

Glu Ser  Gly Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2060                2065                2070

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2075                2080                2085
```

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala
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Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
2105 2110 2115

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
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Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
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Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
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Gly Ser Ala Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu
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Asp Leu Ser Pro Ala Thr Gly His His His His His His
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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
35 40 45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
50 55 60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65 70 75 80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
85 90 95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
100 105 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
115 120 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
130 135 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145 150 155 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

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Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525
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545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800
```

```
Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065
```

```
Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Gly  Ala Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1130                1135                1140

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1145                1150                1155

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1160                1165                1170

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1175                1180                1185

Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1190                1195                1200

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1205                1210                1215

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1220                1225                1230

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1235                1240                1245

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1250                1255                1260

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
    1265                1270                1275

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1280                1285                1290

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1295                1300                1305

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1310                1315                1320
```

```
Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
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Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1340                1345                1350

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1355                1360                1365

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1370                1375                1380

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1385                1390                1395

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1400                1405                1410

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1415                1420                1425

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1430                1435                1440

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1445                1450                1455

Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1460                1465                1470

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1475                1480                1485

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1490                1495                1500

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1505                1510                1515

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1520                1525                1530

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1535                1540                1545

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1550                1555                1560

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1565                1570                1575
```

```
Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1580                1585                1590

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1595                1600                1605

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1610                1615                1620

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1625                1630                1635

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1640                1645                1650

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1655                1660                1665

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1670                1675                1680

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1685                1690                1695

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1700                1705                1710

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1715                1720                1725

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1730                1735                1740

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1745                1750                1755

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1760                1765                1770

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1775                1780                1785

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1790                1795                1800

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1805                1810                1815

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1820                1825                1830
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```
Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1835                 1840                 1845

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    1850                 1855                 1860

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    1865                 1870                 1875

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    1880                 1885                 1890

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1895                 1900                 1905

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    1910                 1915                 1920

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1925                 1930                 1935

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1940                 1945                 1950

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    1955                 1960                 1965

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    1970                 1975                 1980

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1985                 1990                 1995

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2000                 2005                 2010

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser
    2015                 2020                 2025

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Thr Glu
    2030                 2035                 2040

Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2045                 2050                 2055

Glu Ser  Gly Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2060                 2065                 2070

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2075                 2080                 2085
```

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala
2090 2095 2100

Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
2105 2110 2115

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2120 2125 2130

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
2135 2140 2145

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
2150 2155 2160

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
2165 2170 2175

Gly Ser Ala Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu
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Asp Leu Ser Pro Ala Thr Gly His His His His His His
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<210> 1240
<211> 6621
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polypeptide

<400> 1241

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            20                  25                  30

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
        35                  40                  45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
    50                  55                  60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65                  70                  75                  80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
                85                  90                  95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
            100                 105                 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
        115                 120                 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
    130                 135                 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145                 150                 155                 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
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Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
275 280 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
290 295 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305 310 315 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
325 330 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
340 345 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
355 360 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
370 375 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385 390 395 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
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His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
420 425 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
435 440 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
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His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465 470 475 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
485 490 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
500 505 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
515 520 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp

530 535 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545 550 555 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
565 570 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
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Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
595 600 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
610 615 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
645 650 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp

805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
835 840 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
885 890 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
900 905 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
915 920 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
930 935 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945 950 955 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
965 970 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
980 985 990

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
995 1000 1005

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser
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Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
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Ser Ala Pro Gly Ala Ser Ser Pro Pro Val Leu Lys Arg His Gln
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Ala Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
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Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe

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Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln
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Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp
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Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala
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Gln Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
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Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
1145 1150 1155

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
1160 1165 1170

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
1175 1180 1185

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
1190 1195 1200

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
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Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
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Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
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Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
1250 1255 1260

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser
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Ser Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val
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Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg
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Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
1310 1315 1320

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala

1325 1330 1335

Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu
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Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro
1355 1360 1365

Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala
1370 1375 1380

Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser
1385 1390 1395

Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
1400 1405 1410

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg
1415 1420 1425

Gln Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp
1430 1435 1440

Glu Thr Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys
1445 1450 1455

Arg Ala Pro Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu
1460 1465 1470

Asn Tyr Arg Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu
1475 1480 1485

Pro Gly Leu Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu
1490 1495 1500

Leu Ser Met Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser
1505 1510 1515

Gly His Val Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala
1520 1525 1530

Leu Tyr Asn Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu
1535 1540 1545

Pro Ser Lys Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu
1550 1555 1560

His Leu His Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn
1565 1570 1575

Lys Cys Gln Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp

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Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1610                1615                1620

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1625                1630                1635

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1640                1645                1650

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1655                1660                1665

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1670                1675                1680

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1685                1690                1695

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1700                1705                1710

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1715                1720                1725

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1730                1735                1740

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1745                1750                1755

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1760                1765                1770

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1775                1780                1785

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1790                1795                1800

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1805                1810                1815

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1820                1825                1830

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
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1835 1840 1845

Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg
1850 1855 1860

Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu
1865 1870 1875

Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala
1880 1885 1890

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
1895 1900 1905

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
1910 1915 1920

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
1925 1930 1935

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1940 1945 1950

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
1955 1960 1965

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
1970 1975 1980

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
1985 1990 1995

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2000 2005 2010

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser
2015 2020 2025

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu
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Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
2045 2050 2055

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2060 2065 2070

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2075 2080 2085

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala

2090 2095 2100

Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
2105 2110 2115

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2120 2125 2130

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
2135 2140 2145

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
2150 2155 2160

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
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Gly Ser Ala Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu
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Asp Leu Ser Pro Ala Thr Gly His His His His His His
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<212> DNA
<213> Artificial Sequence

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Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
35 40 45

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
50 55 60

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
65 70 75 80

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
85 90 95

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
100 105 110

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
115 120 125

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
130 135 140

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
145 150 155 160

Ser Thr Glu Glu Gly Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
260 265 270

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Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540
```

```
Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815
```

```
Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080
```

```
Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
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Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                 1105                 1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                 1120                 1125

Gln Gly  Ala Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1130                 1135                 1140

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1145                 1150                 1155

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1160                 1165                 1170

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1175                 1180                 1185

Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1190                 1195                 1200

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1205                 1210                 1215

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1220                 1225                 1230

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1235                 1240                 1245

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1250                 1255                 1260

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
    1265                 1270                 1275

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1280                 1285                 1290

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1295                 1300                 1305

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1310                 1315                 1320

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1325                 1330                 1335
```

```
Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
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Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1355                1360                1365

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1370                1375                1380

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1385                1390                1395

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1400                1405                1410

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1415                1420                1425

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1430                1435                1440

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1445                1450                1455

Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1460                1465                1470

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1475                1480                1485

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1490                1495                1500

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1505                1510                1515

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1520                1525                1530

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1535                1540                1545

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1550                1555                1560

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1565                1570                1575

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1580                1585                1590
```

```
Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1595                 1600                 1605

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1610                 1615                 1620

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1625                 1630                 1635

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1640                 1645                 1650

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1655                 1660                 1665

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1670                 1675                 1680

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1685                 1690                 1695

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1700                 1705                 1710

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1715                 1720                 1725

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1730                 1735                 1740

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1745                 1750                 1755

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1760                 1765                 1770

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1775                 1780                 1785

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1790                 1795                 1800

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1805                 1810                 1815

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1820                 1825                 1830

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1835                 1840                 1845
```

```
Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
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Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    1865                1870                1875

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    1880                1885                1890

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1895                1900                1905

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    1910                1915                1920

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1925                1930                1935

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1940                1945                1950

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    1955                1960                1965

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    1970                1975                1980

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1985                1990                1995

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2000                2005                2010

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser
    2015                2020                2025

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Thr Glu
    2030                2035                2040

Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2045                2050                2055

Glu Ser  Gly Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2060                2065                2070

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2075                2080                2085

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Ser  Glu Pro Ala
    2090                2095                2100
```

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Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
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Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
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Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
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Gly Ser Ala Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu
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Asp Leu Ser Pro Ala Thr Gly His His His His His His
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Arg Val Pro Lys Ser Phe Pro Phe Gly Ala Pro Gly Ser Ser Pro Ser
        35                  40                  45

Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
    50                  55                  60

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65                  70                  75                  80

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro
                85                  90                  95

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
            100                 105                 110

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
        115                 120                 125

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser
    130                 135                 140

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145                 150                 155                 160

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
                165                 170                 175

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
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Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
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Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540
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Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
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Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
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```

```
Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
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Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
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Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
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Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
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Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
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Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
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Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
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Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
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Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
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Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
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Gln Gly  Ala Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
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Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
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Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1160                 1165                 1170

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
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Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
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Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
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Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
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Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
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Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
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Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
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Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
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Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
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Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
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Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
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Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
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Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
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Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
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Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
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Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1400                1405                1410

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
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Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1430                1435                1440

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1445                1450                1455

Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
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Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1475                1480                1485

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1490                1495                1500

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1505                1510                1515

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1520                1525                1530

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1535                1540                1545

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1550                1555                1560

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1565                1570                1575

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1580                1585                1590
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Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
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Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
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Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
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Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
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Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
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Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
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Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
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Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
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His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1715                 1720                 1725

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1730                 1735                 1740

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1745                 1750                 1755

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
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Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
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Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
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Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1805                 1810                 1815

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
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Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
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Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
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Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    1865                1870                1875

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
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Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
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Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
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Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1940                1945                1950

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
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Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
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Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
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Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
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Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser
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Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Thr Glu
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Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
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Glu Ser  Gly Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
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Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Ser  Glu Pro Ala
    2090                2095                2100
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Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
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<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1247

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Met Gln Gly Ala Pro Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
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Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
        35                  40                  45

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
    50                  55                  60

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
65                  70                  75                  80

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
                85                  90                  95

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
            100                 105                 110

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
        115                 120                 125

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
    130                 135                 140

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
145                 150                 155                 160

Ser Thr Glu Glu Gly Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270
```

```
Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540
```

```
Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815
```

```
Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080
```

```
Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                 1090                 1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                 1105                 1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                 1120                 1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                 1135                 1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                 1150                 1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                 1165                 1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                 1180                 1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                 1195                 1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                 1210                 1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                 1225                 1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                 1240                 1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                 1255                 1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                 1270                 1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                 1285                 1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1295                 1300                 1305

Pro Ser  Pro Ala Gly Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser
    1310                 1315                 1320

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro
    1325                 1330                 1335
```

```
Ser Glu  Gly Ser Ala Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
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Thr Glu  Glu Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
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Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1370                1375                1380

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
    1385                1390                1395

Ser Gly  Ser Glu Thr Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1400                1405                1410

Glu Thr  Pro Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1415                1420                1425

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1430                1435                1440

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ala Ser  Ser Ala Pro
    1445                1450                1455

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                1555                1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590
```

```
Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                1615                1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845
```

```
Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
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Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                1960                1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                1975                1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                1990                1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                2005                2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                2020                2025

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    2030                2035                2040

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2045                2050                2055

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    2060                2065                2070

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    2075                2080                2085

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    2090                2095                2100
```

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
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Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
2120 2125 2130

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
2135 2140 2145

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2150 2155 2160

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2165 2170 2175

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Ala Thr Gly His His His His His His
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<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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Arg Val Pro Lys Ser Phe Pro Phe Gly Ala Pro Gly Ser Ser Pro Ser
35 40 45

Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
50 55 60

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65 70 75 80

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro
85 90 95

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
100 105 110

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
115 120 125

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser
130 135 140

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145 150 155 160

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
165 170 175

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
260 265 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp

275 280 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
290 295 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305 310 315 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
325 330 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
340 345 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
355 360 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
370 375 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385 390 395 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
405 410 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
420 425 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
435 440 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
450 455 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465 470 475 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
485 490 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
500 505 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
515 520 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
530 535 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln

545 550 555 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
565 570 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
580 585 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
595 600 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
610 615 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
645 650 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met

820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
835 840 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
885 890 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
900 905 910

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915 920 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
930 935 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945 950 955 960

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Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
980 985 990

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
995 1000 1005

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Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
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Ser Ala Pro Gly Ala Ser Ser Pro Pro Val Leu Lys Arg His Gln
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Ala Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
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Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
1070 1075 1080

Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln

1085 1090 1095

Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp
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Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala
1115 1120 1125

Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu
1130 1135 1140

Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
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Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
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Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1175 1180 1185

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1190 1195 1200

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
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Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1220 1225 1230

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1235 1240 1245

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1250 1255 1260

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1265 1270 1275

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1280 1285 1290

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Gly Ala
1295 1300 1305

Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
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Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro
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Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
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Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
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Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr
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Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
1400 1405 1410

Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
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Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
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Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Ala Pro
1445 1450 1455

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1460 1465 1470

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1475 1480 1485

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1490 1495 1500

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
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Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1520 1525 1530

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1535 1540 1545

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
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Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1565 1570 1575

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1580 1585 1590

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg

1595 1600 1605

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1610 1615 1620

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
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Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1640 1645 1650

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1655 1660 1665

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1670 1675 1680

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1685 1690 1695

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
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Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1715 1720 1725

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1730 1735 1740

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
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Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1760 1765 1770

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1775 1780 1785

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1790 1795 1800

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1805 1810 1815

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1820 1825 1830

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1835 1840 1845

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg

1850 1855 1860

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1865 1870 1875

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Pro Gly Thr
1880 1885 1890

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1895 1900 1905

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
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1925 1930 1935

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1940 1945 1950

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
1955 1960 1965

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
1970 1975 1980

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1985 1990 1995

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2000 2005 2010

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
2015 2020 2025

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
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Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2045 2050 2055

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
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Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
2075 2080 2085

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
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Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu

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Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
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Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
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gccacccccg agagtggccc aggcactagt accgagccct ctgaaggcag tgcgccagcc 6540

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Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
        35                  40                  45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
    50                  55                  60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65                  70                  75                  80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
                85                  90                  95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
            100                 105                 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
        115                 120                 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
    130                 135                 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145                 150                 155                 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285
```

```
Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560
```

```
Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830
```

```
Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095
```

```
Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1295                1300                1305

Pro Ser  Pro Ala Gly Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser
    1310                1315                1320

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro
    1325                1330                1335

Ser Glu  Gly Ser Ala Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1340                1345                1350
```

```
Thr Glu  Glu Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
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Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1370                1375                1380

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
    1385                1390                1395

Ser Gly  Ser Glu Thr Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1400                1405                1410

Glu Thr  Pro Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1415                1420                1425

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1430                1435                1440

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ala Ser  Ser Ala Pro
    1445                1450                1455

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                1555                1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605
```

```
Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
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Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860
```

```
Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                1960                1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                1975                1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                1990                1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                2005                2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                2020                2025

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    2030                2035                2040

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2045                2050                2055

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    2060                2065                2070

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    2075                2080                2085

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    2090                2095                2100

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
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Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
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Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
2135 2140 2145

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2150 2155 2160

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2165 2170 2175

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Ala Thr Gly His His His His His His
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Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
        35                  40                  45

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
    50                  55                  60

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
65                  70                  75                  80

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
                85                  90                  95

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
            100                 105                 110

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
        115                 120                 125

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
    130                 135                 140

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
145                 150                 155                 160

Ser Thr Glu Glu Gly Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285
```

```
Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560
```

```
Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830
```

```
Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095
```

```
Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
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Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1295                1300                1305

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala
    1310                1315                1320

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1325                1330                1335

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1340                1345                1350
```

```
Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
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Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala
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Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1385                 1390                 1395

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1400                 1405                 1410

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1415                 1420                 1425

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ala
    1430                 1435                 1440

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Ala Ser  Ser Ala Pro
    1445                 1450                 1455

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                 1465                 1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                 1480                 1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                 1495                 1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                 1510                 1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                 1525                 1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                 1540                 1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                 1555                 1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                 1570                 1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                 1585                 1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                 1600                 1605
```

```
Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
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Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860
```

```
Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
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Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                1960                1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                1975                1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                1990                1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                2005                2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                2020                2025

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    2030                2035                2040

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2045                2050                2055

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    2060                2065                2070

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    2075                2080                2085

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    2090                2095                2100

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
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Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
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Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
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Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
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Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Ala Thr Gly His His His His His His
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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
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Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
    50                  55                  60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65                  70                  75                  80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
                85                  90                  95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
            100                 105                 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
        115                 120                 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
    130                 135                 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145                 150                 155                 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285
```

```
Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560
```

```
Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830
```

```
Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095
```

```
Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
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Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1295                1300                1305

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala
    1310                1315                1320

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1325                1330                1335

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1340                1345                1350
```

```
Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
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Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala
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Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1385                 1390                 1395

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1400                 1405                 1410

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1415                 1420                 1425

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ala
    1430                 1435                 1440

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Ala Ser  Ser Ala Pro
    1445                 1450                 1455

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                 1465                 1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                 1480                 1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                 1495                 1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                 1510                 1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                 1525                 1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                 1540                 1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                 1555                 1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                 1570                 1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                 1585                 1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                 1600                 1605
```

```
Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
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Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860
```

```
Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                1960                1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                1975                1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                1990                1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                2005                2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                2020                2025

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    2030                2035                2040

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2045                2050                2055

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    2060                2065                2070

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    2075                2080                2085

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
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Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
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Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
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Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
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Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
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Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65                  70                  75                  80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
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Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
            100                 105                 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
        115                 120                 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
    130                 135                 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145                 150                 155                 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
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Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
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Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
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Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
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Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
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Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
325 330 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
340 345 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
355 360 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
370 375 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385 390 395 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
405 410 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
420 425 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
435 440 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
450 455 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465 470 475 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
485 490 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
500 505 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
515 520 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
530 535 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545 550 555 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val

565 570 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
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Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
595 600 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
610 615 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
645 650 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg

835 840 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
885 890 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
900 905 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
915 920 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
930 935 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
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980 985 990

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
995 1000 1005

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser
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Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
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Ser Ala Pro Gly Ala Ser Ser Pro Pro Val Leu Lys Arg His Gln
1040 1045 1050

Ala Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
1055 1060 1065

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
1070 1075 1080

Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln
1085 1090 1095

Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp

1100 1105 1110

Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala
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Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu
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Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
1145 1150 1155

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1160 1165 1170

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1175 1180 1185

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1190 1195 1200

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1205 1210 1215

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1220 1225 1230

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1235 1240 1245

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1250 1255 1260

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1265 1270 1275

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1280 1285 1290

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Gly Ala
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Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly

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1370 1375 1380

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
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Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
1400 1405 1410

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
1415 1420 1425

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala
1430 1435 1440

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Ala Pro
1445 1450 1455

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1460 1465 1470

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1475 1480 1485

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1490 1495 1500

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
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Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
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1535 1540 1545

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1550 1555 1560

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1565 1570 1575

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1580 1585 1590

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1595 1600 1605

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro

1610 1615 1620

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
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Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1640 1645 1650

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1655 1660 1665

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1670 1675 1680

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1685 1690 1695

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
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1715 1720 1725

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1730 1735 1740

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1745 1750 1755

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1760 1765 1770

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1775 1780 1785

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1805 1810 1815

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Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1835 1840 1845

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1865 1870 1875

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1880 1885 1890

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1895 1900 1905

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1940 1945 1950

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
1955 1960 1965

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1970 1975 1980

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1985 1990 1995

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2000 2005 2010

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
2015 2020 2025

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
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Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
    50                  55                  60

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65                  70                  75                  80

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro
                85                  90                  95

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
            100                 105                 110

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
        115                 120                 125

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser
    130                 135                 140

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145                 150                 155                 160

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
                165                 170                 175

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300
```

```
Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
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Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575
```

```
Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845
```

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Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110
```

```
Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
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Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1295                1300                1305

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala
    1310                1315                1320

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1325                1330                1335

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
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Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
    1355                1360                1365
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Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala
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Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
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Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1400                1405                1410

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
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Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ala
    1430                1435                1440

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Ala Ser  Ser Ala Pro
    1445                1450                1455

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                1555                1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                1615                1620
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Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875
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Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                1960                1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                1975                1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                1990                1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                2005                2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                2020                2025

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    2030                2035                2040

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    2060                2065                2070

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    2075                2080                2085

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    2090                2095                2100

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    2105                2110                2115

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    2120                2125                2130
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Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
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Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
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Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

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Arg Val Pro Lys Ser Phe Pro Phe Gly Ala Pro Gly Ser Ser Pro Ser
        35                  40                  45

Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
    50                  55                  60

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65                  70                  75                  80

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro
                85                  90                  95

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
            100                 105                 110

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
        115                 120                 125

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser
    130                 135                 140

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145                 150                 155                 160

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
                165                 170                 175

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300
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Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575
```

```
Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845
```

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Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110
```

```
Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gly Ala Pro Ser  Pro Ala Gly Ser Pro  Thr Ser Thr
    1310                1315                1320

Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    1325                1330                1335

Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala Pro Gly  Ser Pro Ala
    1340                1345                1350

Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr Ser Thr  Glu Pro Ser
    1355                1360                1365
```

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Glu Gly  Ser Ala Pro Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    1370                1375                1380

Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    1385                1390                1395

Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr Pro Gly  Ser Glu Pro
    1400                1405                1410

Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Ser Pro Ala  Gly Ser Pro
    1415                1420                1425

Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1430                1435                1440

Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    1445                1450                1455

Ala Ser  Ser Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                1555                1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                1615                1620
```

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Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875
```

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Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                1960                1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                1975                1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                1990                1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                2005                2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                2020                2025

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    2030                2035                2040

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2045                2050                2055

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    2060                2065                2070

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    2075                2080                2085

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    2090                2095                2100

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    2105                2110                2115

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    2120                2125                2130
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Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
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Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2150 2155 2160

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Ala Thr Gly His His His His His His
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Gly Ser Ser
20 25 30

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        35                  40                  45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
    50                  55                  60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65                  70                  75                  80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
                85                  90                  95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
            100                 105                 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
        115                 120                 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
    130                 135                 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145                 150                 155                 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300
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Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575
```

```
Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845
```

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Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110
```

```
Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                 1120                 1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                 1135                 1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                 1150                 1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                 1165                 1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                 1180                 1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                 1195                 1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                 1210                 1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                 1225                 1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                 1240                 1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                 1255                 1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                 1270                 1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                 1285                 1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                 1300                 1305

Cys Asn  Ile Gly Ala Pro Ser  Pro Ala Gly Ser Pro  Thr Ser Thr
    1310                 1315                 1320

Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    1325                 1330                 1335

Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala Pro Gly  Ser Pro Ala
    1340                 1345                 1350

Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr Ser Thr  Glu Pro Ser
    1355                 1360                 1365
```

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Glu Gly  Ser Ala Pro Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    1370                1375                1380

Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    1385                1390                1395

Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr Pro Gly  Ser Glu Pro
    1400                1405                1410

Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Ser Pro Ala  Gly Ser Pro
    1415                1420                1425

Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1430                1435                1440

Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    1445                1450                1455

Ala Ser  Ser Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                1555                1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                1615                1620
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Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875
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Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                1960                1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                1975                1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                1990                1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                2005                2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                2020                2025

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    2030                2035                2040

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2045                2050                2055

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    2060                2065                2070

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    2075                2080                2085

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    2090                2095                2100

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    2105                2110                2115

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    2120                2125                2130
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Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
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Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2150 2155 2160

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Ala Thr Gly His His His His His His
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<212> PRT
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Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu

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Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
    50                  55                  60

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
65                  70                  75                  80

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
                85                  90                  95

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
            100                 105                 110

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
        115                 120                 125

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
    130                 135                 140

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
145                 150                 155                 160

Ser Thr Glu Glu Gly Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
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305 310 315 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
325 330 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
340 345 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
355 360 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
370 375 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385 390 395 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
405 410 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
420 425 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
435 440 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
450 455 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465 470 475 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
485 490 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
500 505 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
515 520 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
530 535 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545 550 555 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
565 570 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile

580 585 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
595 600 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
610 615 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
645 650 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
835 840 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn

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Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
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Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu
1130 1135 1140

Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
1145 1150 1155

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1160 1165 1170

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1175 1180 1185

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1190 1195 1200

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1205 1210 1215

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1220 1225 1230

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1235 1240 1245

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1250 1255 1260

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1265 1270 1275

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1280 1285 1290

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1295 1300 1305

Cys Asn Ile Gly Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr
1310 1315 1320

Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
1325 1330 1335

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
1340 1345 1350

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
1355 1360 1365

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser

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Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    1385                1390                1395

Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr Pro Gly  Ser Glu Pro
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Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Ser Pro Ala  Gly Ser Pro
    1415                1420                1425

Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1430                1435                1440

Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    1445                1450                1455

Ala Ser  Ser Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                1555                1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                1615                1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
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Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1640 1645 1650

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1655 1660 1665

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1670 1675 1680

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1685 1690 1695

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1700 1705 1710

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1715 1720 1725

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1730 1735 1740

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1745 1750 1755

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1760 1765 1770

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
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Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1790 1795 1800

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1805 1810 1815

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1820 1825 1830

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1835 1840 1845

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1850 1855 1860

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1865 1870 1875

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Pro Gly Thr

1880 1885 1890

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1895 1900 1905

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1910 1915 1920

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1925 1930 1935

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1940 1945 1950

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
1955 1960 1965

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
1970 1975 1980

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1985 1990 1995

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2000 2005 2010

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
2015 2020 2025

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
2030 2035 2040

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2045 2050 2055

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
2060 2065 2070

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
2075 2080 2085

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
2090 2095 2100

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
2105 2110 2115

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
2120 2125 2130

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala

2135 2140 2145

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2150 2155 2160

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Ala Thr Gly His His His His His His
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tcggaaacgc caggcacaag tgagtctgca actcccgagt ccggacctgg ctccgagcct 180

gccactagcg gctccgagac tccgggaact tccgagagcg ctacaccaga aagcggaccc 240

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
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Arg Val Pro Lys Ser Phe Pro Phe Gly Ala Pro Gly Ser Ser Pro Ser
35 40 45

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Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
    50                  55                  60

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65                  70                  75                  80

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro
                85                  90                  95

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
            100                 105                 110

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
        115                 120                 125

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser
    130                 135                 140

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145                 150                 155                 160

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
                165                 170                 175

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320
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Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590
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Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860
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Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125
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Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
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Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gly Ala Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    1310                1315                1320

Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1325                1330                1335

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
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Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala
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Ser Thr  Gly Thr Gly Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
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Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
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Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
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Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser
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Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
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Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1445                 1450                 1455

Ala Ser  Ser Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
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Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                 1480                 1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                 1495                 1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                 1510                 1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                 1525                 1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                 1540                 1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                 1555                 1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
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Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
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Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                 1600                 1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                 1615                 1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
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Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
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Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
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Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
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Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
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Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
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Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
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Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
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Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
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Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
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Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
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Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
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Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
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Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
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Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
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Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                1990                1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
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Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
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Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
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Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
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Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
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Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
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Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
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Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Glu Pro Ala
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Pro Ala Thr Gly His His His His His His
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<223> Description of Artificial Sequence: Synthetic polynucleotide

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<212> PRT
<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Gly Ser Ser
20 25 30

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
35 40 45

```
Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
    50                  55                  60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65                  70                  75                  80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
                85                  90                  95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
            100                 105                 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
        115                 120                 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
    130                 135                 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145                 150                 155                 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320
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Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590
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```
Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860
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Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125
```

```
Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gly Ala Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    1310                1315                1320

Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1325                1330                1335

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1340                1345                1350

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala
    1355                1360                1365

Ser Thr  Gly Thr Gly Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    1370                1375                1380
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Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1385                1390                1395

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
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Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser
    1415                1420                1425

Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    1430                1435                1440

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1445                1450                1455

Ala Ser  Ser Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                1555                1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                1615                1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635
```

```
Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                1960                1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                1975                1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                1990                1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                2005                2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                2020                2025

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    2030                2035                2040

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2045                2050                2055

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    2060                2065                2070

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    2075                2080                2085

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    2090                2095                2100

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    2105                2110                2115

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    2120                2125                2130

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Glu Pro Ala
    2135                2140                2145
```

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2150 2155 2160

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2165 2170 2175

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Ala Thr Gly His His His His His His
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<210> 1270
<211> 6612
<212> DNA
<213> Artificial Sequence

<220>
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ccaggttcgt caccctcagc atcaacgggg acgggaccgg gggcgtcacc aggaacgtcc 180

tccaccggct cgccgggtgc atcacccgga acgtcatcga ccggatcgcc agggagctcg 240

acgccatcag gcgcaacagg atcacctggc tcaagcccta gcgcgtcaac cggcacgggt 300

ccgggtgcct cccctggcac gtccagcacc ggatcacccg gatcgagccc atccgcctca 360

accggaaccg gacccggtac accagggtcg ggaacagcct cctcgtcacc aggctcctca 420

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cccggagcat cgccaggaac ttcgagcacg gggtcgcccg cctcgagcga cgcaagattt 540

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ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

ggcctcattg gagccctact agtatgtaga gaagggagtc tggccaagga aaagacacag 1020

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<210> 1271
<211> 2203
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Gly Ala Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
20 25 30

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
35 40 45

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Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
    50                  55                  60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65                  70                  75                  80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
                85                  90                  95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
            100                 105                 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
        115                 120                 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
    130                 135                 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145                 150                 155                 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320
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Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590
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Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860
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Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125
```

```
Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gly Ala Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    1310                1315                1320

Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1325                1330                1335

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1340                1345                1350

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala
    1355                1360                1365

Ser Thr  Gly Thr Gly Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    1370                1375                1380
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Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1385                1390                1395

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
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Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser
    1415                1420                1425

Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    1430                1435                1440

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1445                1450                1455

Ala Ser  Ser Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                1555                1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                1615                1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635
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Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                1960                1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                1975                1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                1990                1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                2005                2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                2020                2025

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    2030                2035                2040

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2045                2050                2055

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    2060                2065                2070

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    2075                2080                2085

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    2090                2095                2100

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    2105                2110                2115

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    2120                2125                2130

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Glu Pro Ala
    2135                2140                2145
```

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2150 2155 2160

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2165 2170 2175

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
2180 2185 2190

Pro Ala Thr Gly His His His His His His
2195 2200

<210> 1272
<211> 6612
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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ccaggctcca gcccctccgc gagcacggga accggaccag gttcgtcacc ctcagcatca 120

acggggacgg gaccgggggc gtcaccagga acgtcctcca ccggctcgcc gggtgcatca 180

cccggaacgt catcgaccgg atcgccaggg agctcgacgc catcaggcgc aacaggatca 240

cctggctcaa gccctagcgc gtcaaccggc acgggtccgg gtgcctcccc tggcacgtcc 300

agcaccggat cacccggatc gagcccatcc gcctcaaccg gaaccggacc cggtacacca 360

gggtcgggaa cagcctcctc gtcaccaggc tcctcaaccc cctcgggagc cacgggttcg 420

cccggttcgt caacgccttc cggagcaact ggtagccccg gagcatcgcc aggaacttcg 480

agcacggggt cgcccgcctc gagcagtgat ctcggtgagc tgcctgtgga cgcaagattt 540

cctcctagag tgccaaaatc ttttccattc aacacctcag tcgtgtacaa aaagactctg 600

tttgtagaat tcacggatca ccttttcaac atcgctaagc caaggccacc ctggatgggt 660

ctgctaggtc ctaccatcca ggctgaggtt tatgatacag tggtcattac acttaagaac 720

atggcttccc atcctgtcag tcttcatgct gttggtgtat cctactggaa agcttctgag 780

ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct 840

ggtggaagcc atacatatgt ctggcaggtc ctgaaagaga atggtccaat ggcctctgac 900

ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

ggcctcattg gagccctact agtatgtaga gaagggagtc tggccaagga aaagacacag 1020

accttgcaca aatttatact acttttgct gtatttgatg aagggaaaag ttggcactca 1080

gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg 1200

aaatcagtct attggcatgt gattggaatg ggcaccactc ctgaagtgca ctcaatattc 1260

ctcgaaggtc acacatttct tgtgaggaac catcgccagg ctagcttgga aatctcgcca 1320

ataactttcc ttactgctca aacactcttg atggaccttg gacagtttct actgttttgt 1380

catatctctt cccaccaaca tgatggcatg gaagcttatg tcaaagtaga cagctgtcca 1440

gaggaacccc aactacgaat gaaaaataat gaagaagcgg aagactatga tgatgatctt 1500

actgattctg aaatggatgt ggtcaggttt gatgatgaca actctccttc ctttatccaa 1560

attcgctcag ttgccaagaa gcatcctaaa acttgggtac attacattgc tgctgaagag 1620

gaggactggg actatgctcc cttagtcctc gcccccgatg acagaagtta taaaagtcaa 1680

tatttgaaca atggccctca gcggattggt aggaagtaca aaaaagtccg atttatggca 1740

tacacagatg aaacctttaa gactcgtgaa gctattcagc atgaatcagg aatcttggga 1800

cctttacttt atggggaagt tggagacaca ctgttgatta tatttaagaa tcaagcaagc 1860

agaccatata acatctaccc tcacggaatc actgatgtcc gtcctttgta ttcaaggaga 1920

ttaccaaaag gtgtaaaaca tttgaaggat tttccaattc tgccaggaga aatattcaaa 1980

tataaatgga cagtgactgt agaagatggg ccaactaaat cagatcctcg gtgcctgacc 2040

cgctattact ctagtttcgt taatatggag agagatctag cttcaggact cattggccct 2100

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catttgggac tcctggggcc atatataaga gcagaagttg aagataatat catggtaact 3540

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tgggcttatt tctctgatgt tgacctggaa aaagatgtgc actcaggcct gattggaccc 3780

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acctctgagt ccgccacacc agagtccgga cccggatctc ccgctgggag ccccacctcc 6060

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ccatccgagg gatcagctcc cggcacctca gagtcggcaa ccccggagtc tggacccgga 6180

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tccggccctg ggtctgaacc cgccacaagt ggtagtgaga caccaggatc agaacctgct 6300

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tcagcacccg gctctgagcc ggccacaagt ggcagtgaga cacccggcac ttcagagagt 6480

gccacccccg agagtggccc aggcactagt accgagccct ctgaaggcag tgcgccagcc 6540

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catcaccatt ga 6612

<210> 1273
<211> 2203
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1273
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Met Gln Gly Ala Pro Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
20 25 30

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
35 40 45

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly

50 55 60

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
65 70 75 80

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
85 90 95

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
100 105 110

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
115 120 125

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
130 135 140

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
145 150 155 160

Ser Thr Glu Glu Gly Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
260 265 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
275 280 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
290 295 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305 310 315 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys

325 330 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
340 345 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
355 360 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
370 375 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385 390 395 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
405 410 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
420 425 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
435 440 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
450 455 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465 470 475 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
485 490 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
500 505 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
515 520 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
530 535 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545 550 555 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
565 570 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
580 585 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly

595 600 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
610 615 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
645 650 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
835 840 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu

865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
885 890 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
900 905 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
915 920 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
930 935 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945 950 955 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
965 970 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
980 985 990

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
995 1000 1005

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser
1010 1015 1020

Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
1025 1030 1035

Ser Ala Pro Gly Ala Ser Ser Pro Pro Val Leu Lys Arg His Gln
1040 1045 1050

Ala Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
1055 1060 1065

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
1070 1075 1080

Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln
1085 1090 1095

Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp
1100 1105 1110

Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala
1115 1120 1125

Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu

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Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gly Ala Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    1310                1315                1320

Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1325                1330                1335

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1340                1345                1350

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala
    1355                1360                1365

Ser Thr  Gly Thr Gly Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    1370                1375                1380

Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
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1385 1390 1395

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
1400 1405 1410

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser
1415 1420 1425

Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
1430 1435 1440

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
1445 1450 1455

Ala Ser Ser Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1460 1465 1470

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1475 1480 1485

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1490 1495 1500

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1505 1510 1515

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1520 1525 1530

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1535 1540 1545

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1550 1555 1560

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1565 1570 1575

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1580 1585 1590

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1595 1600 1605

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1610 1615 1620

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1625 1630 1635

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr

1640 1645 1650

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1655 1660 1665

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1670 1675 1680

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1685 1690 1695

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1700 1705 1710

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1715 1720 1725

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1730 1735 1740

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1745 1750 1755

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1760 1765 1770

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1775 1780 1785

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1790 1795 1800

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1805 1810 1815

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1820 1825 1830

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1835 1840 1845

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1850 1855 1860

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1865 1870 1875

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Pro Gly Thr
1880 1885 1890

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala

1895 1900 1905

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
1910 1915 1920

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1925 1930 1935

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1940 1945 1950

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
1955 1960 1965

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
1970 1975 1980

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1985 1990 1995

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2000 2005 2010

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
2015 2020 2025

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
2030 2035 2040

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2045 2050 2055

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
2060 2065 2070

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
2075 2080 2085

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
2090 2095 2100

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
2105 2110 2115

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
2120 2125 2130

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
2135 2140 2145

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro

2150 2155 2160

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2165 2170 2175

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
2180 2185 2190

Pro Ala Thr Gly His His His His His His
2195 2200

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<211> 6612
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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tcggaaacgc caggcacaag tgagtctgca actcccgagt ccggacctgg ctccgagcct 180

gccactagcg gctccgagac tccgggaact tccgagagcg ctacaccaga aagcggaccc 240

ggaaccagta ccgaacctag cgagggctct gctccgggca gcccagccgg ctctcctaca 300

tccacggagg agggcacttc cgaatccgcc accccggagt cagggccagg atctgaaccc 360

gctacctcag gcagtgagac gccaggaacg agcgagtccg ctacaccgga gagtgggcca 420

gggagccctg ctggatctcc tacgtccact gaggaagggt caccagcggg ctcgcccacc 480

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ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

ggcctcattg gagccctact agtatgtaga gaagggagtc tggccaagga aaagacacag 1020

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gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

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tcgagcgaac aaaaactcat ctcagaagag gatctgtcac ctgcaaccgg tcatcatcac 6600

catcaccatt ga 6612

<210> 1275
<211> 2203
<212> PRT
<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Gly Ala Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
20 25 30

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
35 40 45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
50 55 60

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Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65                  70                  75                  80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
                85                  90                  95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
            100                 105                 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
        115                 120                 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
    130                 135                 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145                 150                 155                 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335
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Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605
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Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880
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Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140
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```
Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Gly Ala Pro Ser  Pro Ala Gly
    1310                1315                1320

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1325                1330                1335

Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    1340                1345                1350

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr
    1355                1360                1365

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Thr  Ser Thr Glu
    1370                1375                1380

Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1385                1390                1395
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Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
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Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Ser
    1415                 1420                 1425

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    1430                 1435                 1440

Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu
    1445                 1450                 1455

Gly Ser  Ala Pro Gly Ala Ser  Ser Thr Phe Lys Glu  Asn Tyr Arg
    1460                 1465                 1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                 1480                 1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                 1495                 1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                 1510                 1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                 1525                 1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                 1540                 1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                 1555                 1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                 1570                 1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                 1585                 1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                 1600                 1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                 1615                 1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                 1630                 1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                 1645                 1650
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```
Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905
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```
Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                 1915                 1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                 1930                 1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                 1945                 1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                 1960                 1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                 1975                 1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                 1990                 1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                 2005                 2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                 2020                 2025

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    2030                 2035                 2040

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2045                 2050                 2055

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    2060                 2065                 2070

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    2075                 2080                 2085

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    2090                 2095                 2100

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    2105                 2110                 2115

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    2120                 2125                 2130

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Glu Pro Ala
    2135                 2140                 2145

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2150                 2155                 2160
```

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2165 2170 2175

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
2180 2185 2190

Pro Ala Thr Gly His His His His His His
2195 2200

<210> 1276
<211> 6612
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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acggggacgg gaccgggggc gtcaccagga acgtcctcca ccggctcgcc gggtgcatca 180

cccggaacgt catcgaccgg atcgccaggg agctcgacgc catcaggcgc aacaggatca 240

cctggctcaa gccctagcgc gtcaaccggc acgggtccgg gtgcctcccc tggcacgtcc 300

agcaccggat cacccggatc gagcccatcc gcctcaaccg gaaccggacc cggtacacca 360

gggtcgggaa cagcctcctc gtcaccaggc tcctcaaccc cctcgggagc cacgggttcg 420

cccggttcgt caacgccttc cggagcaact ggtagccccg gagcatcgcc aggaacttcg 480

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ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

ggcctcattg gagccctact agtatgtaga gaagggagtc tggccaagga aaagacacag 1020

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gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

atgcacacag tcaatggtta tgtaaacagg tctctgccag gtctgattgg atgccacagg 1200

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cctttacttt atggggaagt tggagacaca ctgttgatta tatttaagaa tcaagcaagc 1860

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acctcagggt cagagacacc cggatctccg gcaggctcac caacctccac tgaggagggc 6360

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tcagcacccg gctctgagcc ggccacaagt ggcagtgaga cacccggcac ttcagagagt 6480

gccacccccg agagtggccc aggcactagt accgagccct ctgaaggcag tgcgccagcc 6540

tcgagcgaac aaaaactcat ctcagaagag gatctgtcac ctgcaaccgg tcatcatcac 6600

catcaccatt ga 6612

<210> 1277
<211> 2203
<212> PRT
<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Gly Ala Pro Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
20 25 30

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
35 40 45

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
50 55 60

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Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
65                  70                  75                  80

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
                85                  90                  95

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
            100                 105                 110

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
        115                 120                 125

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
    130                 135                 140

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
145                 150                 155                 160

Ser Thr Glu Glu Gly Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335
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Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605
```

```
Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880
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Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140
```

```
Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Gly Ala Pro Ser  Pro Ala Gly
    1310                1315                1320

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1325                1330                1335

Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    1340                1345                1350

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr
    1355                1360                1365

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Thr  Ser Thr Glu
    1370                1375                1380

Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1385                1390                1395
```

```
Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
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Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Ser
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Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
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Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu
    1445                 1450                 1455

Gly Ser  Ala Pro Gly Ala Ser  Ser Thr Phe Lys Glu  Asn Tyr Arg
    1460                 1465                 1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                 1480                 1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                 1495                 1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                 1510                 1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                 1525                 1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                 1540                 1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                 1555                 1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                 1570                 1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                 1585                 1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                 1600                 1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                 1615                 1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                 1630                 1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
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```
Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
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Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905
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Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                 1915                 1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                 1930                 1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                 1945                 1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                 1960                 1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                 1975                 1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                 1990                 1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                 2005                 2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                 2020                 2025

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    2030                 2035                 2040

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2045                 2050                 2055

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    2060                 2065                 2070

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    2075                 2080                 2085

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    2090                 2095                 2100

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    2105                 2110                 2115

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    2120                 2125                 2130

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Glu Pro Ala
    2135                 2140                 2145

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2150                 2155                 2160
```

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2165 2170 2175

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
2180 2185 2190

Pro Ala Thr Gly His His His His His His
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<210> 1278
<211> 6612
<212> DNA
<213> Artificial Sequence

<220>
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gccactagcg gctccgagac tccgggaact tccgagagcg ctacaccaga aagcggaccc 240

ggaaccagta ccgaacctag cgagggctct gctccgggca gcccagccgg ctctcctaca 300

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gctacctcag gcagtgagac gccaggaacg agcgagtccg ctacaccgga gagtgggcca 420

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catcaccatt ga 6612

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<211> 2203
<212> PRT
<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Gly Ala Pro Gly Ser Ser Pro Ser
35 40 45

Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
50 55 60

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Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65                  70                  75                  80

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro
                85                  90                  95

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
            100                 105                 110

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
        115                 120                 125

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser
    130                 135                 140

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145                 150                 155                 160

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
                165                 170                 175

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335
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Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605
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Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880
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Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140
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Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
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Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
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Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Gly Ala Pro Ser  Pro Ala Gly
    1310                1315                1320

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
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Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
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Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr
    1355                1360                1365

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Thr  Ser Thr Glu
    1370                1375                1380

Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1385                1390                1395
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Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
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Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Ser
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Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    1430                 1435                 1440

Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu
    1445                 1450                 1455

Gly Ser  Ala Pro Gly Ala Ser  Ser Thr Phe Lys Glu  Asn Tyr Arg
    1460                 1465                 1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                 1480                 1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                 1495                 1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                 1510                 1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                 1525                 1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                 1540                 1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                 1555                 1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                 1570                 1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                 1585                 1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                 1600                 1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                 1615                 1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                 1630                 1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                 1645                 1650
```

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Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
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Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905
```

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
1910 1915 1920

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1925 1930 1935

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1940 1945 1950

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
1955 1960 1965

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
1970 1975 1980

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1985 1990 1995

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2000 2005 2010

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
2015 2020 2025

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
2030 2035 2040

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2045 2050 2055

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
2060 2065 2070

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
2075 2080 2085

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
2090 2095 2100

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
2105 2110 2115

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
2120 2125 2130

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
2135 2140 2145

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2150 2155 2160

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2165 2170 2175

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
2180 2185 2190

Pro Ala Thr Gly His His His His His His
2195 2200

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<211> 6612
<212> DNA
<213> Artificial Sequence

<220>
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gcatcaacgg ggacgggacc gggggcgtca ccaggaacgt cctccaccgg ctcgccgggt 240

gcatcacccg gaacgtcatc gaccggatcg ccagggagct cgacgccatc aggcgcaaca 300

ggatcacctg gctcaagccc tagcgcgtca accggcacgg gtccgggtgc ctcccctggc 360

acgtccagca ccggatcacc cggatcgagc ccatccgcct caaccggaac cggacccggt 420

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ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

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catcaccatt ga 6612

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<212> PRT
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Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
35 40 45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
50 55 60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser

65 70 75 80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
85 90 95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
100 105 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
115 120 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
130 135 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145 150 155 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
260 265 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
275 280 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
290 295 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305 310 315 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
325 330 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe

340 345 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
355 360 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
370 375 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385 390 395 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
405 410 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
420 425 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
435 440 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
450 455 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465 470 475 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
485 490 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
500 505 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
515 520 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
530 535 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545 550 555 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
565 570 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
580 585 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
595 600 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn

610 615 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
645 650 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
835 840 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly

885 890 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
900 905 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
915 920 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
930 935 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945 950 955 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
965 970 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
980 985 990

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
995 1000 1005

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1010 1015 1020

Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
1025 1030 1035

Ser Ala Pro Gly Ala Ser Ser Pro Pro Val Leu Lys Arg His Gln
1040 1045 1050

Ala Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
1055 1060 1065

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
1070 1075 1080

Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln
1085 1090 1095

Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp
1100 1105 1110

Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala
1115 1120 1125

Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu
1130 1135 1140

Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu

1145 1150 1155

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1160 1165 1170

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1175 1180 1185

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1190 1195 1200

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1205 1210 1215

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1220 1225 1230

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1235 1240 1245

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1250 1255 1260

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1265 1270 1275

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1280 1285 1290

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1295 1300 1305

Cys Asn Ile Gln Met Glu Asp Pro Gly Ala Pro Ser Pro Ala Gly
1310 1315 1320

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
1325 1330 1335

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
1340 1345 1350

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr
1355 1360 1365

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
1370 1375 1380

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
1385 1390 1395

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr

1400 1405 1410

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser
1415 1420 1425

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
1430 1435 1440

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
1445 1450 1455

Gly Ser Ala Pro Gly Ala Ser Ser Thr Phe Lys Glu Asn Tyr Arg
1460 1465 1470

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1475 1480 1485

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1490 1495 1500

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1505 1510 1515

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1520 1525 1530

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1535 1540 1545

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1550 1555 1560

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1565 1570 1575

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1580 1585 1590

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1595 1600 1605

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1610 1615 1620

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1625 1630 1635

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1640 1645 1650

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp

1655 1660 1665

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1670 1675 1680

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1685 1690 1695

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1700 1705 1710

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1715 1720 1725

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1730 1735 1740

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1745 1750 1755

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1760 1765 1770

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1775 1780 1785

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1790 1795 1800

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1805 1810 1815

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1820 1825 1830

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1835 1840 1845

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1850 1855 1860

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1865 1870 1875

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Pro Gly Thr
1880 1885 1890

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1895 1900 1905

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro

1910 1915 1920

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1925 1930 1935

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1940 1945 1950

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
1955 1960 1965

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
1970 1975 1980

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1985 1990 1995

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2000 2005 2010

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
2015 2020 2025

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
2030 2035 2040

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2045 2050 2055

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
2060 2065 2070

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
2075 2080 2085

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
2090 2095 2100

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
2105 2110 2115

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
2120 2125 2130

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
2135 2140 2145

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2150 2155 2160

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala

2165 2170 2175

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
2180 2185 2190

Pro Ala Thr Gly His His His His His His
2195 2200

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<211> 6612
<212> DNA
<213> Artificial Sequence

<220>
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ccaggttcgt caccctcagc atcaacgggg acgggaccgg gggcgtcacc aggaacgtcc 180

tccaccggct cgccgggtgc atcacccgga acgtcatcga ccggatcgcc agggagctcg 240

acgccatcag gcgcaacagg atcacctggc tcaagcccta gcgcgtcaac cggcacgggt 300

ccgggtgcct cccctggcac gtccagcacc ggatcacccg gatcgagccc atccgcctca 360

accggaaccg gacccggtac accagggtcg ggaacagcct cctcgtcacc aggctcctca 420

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cccggagcat cgccaggaac ttcgagcacg gggtcgcccg cctcgagcga cgcaagattt 540

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ggagctgaat atgatgatca gaccagtcaa agggagaaag aagatgataa agtcttccct 840

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ccactgtgcc ttacctactc atatctttct catgtggacc tggtaaaaga cttgaattca 960

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tcagcacccg gctctgagcc ggccacaagt ggcagtgaga cacccggcac ttcagagagt 6480

gccacccccg agagtggccc aggcactagt accgagccct ctgaaggcag tgcgccagcc 6540

tcgagcgaac aaaaactcat ctcagaagag gatctgtcac ctgcaaccgg tcatcatcac 6600

catcaccatt ga 6612

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<211> 2203
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Gly Ala Pro Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
20 25 30

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
35 40 45

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
50 55 60

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
65 70 75 80

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Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
                85                  90                  95

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
            100                 105                 110

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
        115                 120                 125

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
    130                 135                 140

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
145                 150                 155                 160

Ser Thr Glu Glu Gly Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350
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Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620
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Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895
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Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155
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```
Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
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Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Gly Ala Pro Gly  Thr Pro Gly
    1310                1315                1320

Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1325                1330                1335

Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1340                1345                1350

Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly
    1355                1360                1365

Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly Pro Gly  Thr Pro Gly
    1370                1375                1380

Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1385                1390                1395

Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1400                1405                1410
```

```
Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly
    1415                 1420                 1425

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr
    1430                 1435                 1440

Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1445                 1450                 1455

Ser Thr  Gly Ser Pro Ala Ser  Ser Thr Phe Lys Glu  Asn Tyr Arg
    1460                 1465                 1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                 1480                 1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                 1495                 1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                 1510                 1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                 1525                 1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                 1540                 1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                 1555                 1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                 1570                 1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                 1585                 1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                 1600                 1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                 1615                 1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                 1630                 1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                 1645                 1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                 1660                 1665
```

```
Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920
```

```
Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                1960                1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                1975                1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                1990                1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                2005                2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                2020                2025

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    2030                2035                2040

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2045                2050                2055

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    2060                2065                2070

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    2075                2080                2085

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    2090                2095                2100

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    2105                2110                2115

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    2120                2125                2130

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Glu Pro Ala
    2135                2140                2145

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2150                2155                2160

Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    2165                2170                2175
```

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
2180 2185 2190

Pro Ala Thr Gly His His His His His His
2195 2200

<210> 1284
<211> 6612
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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tcggaaacgc caggcacaag tgagtctgca actcccgagt ccggacctgg ctccgagcct 180

gccactagcg gctccgagac tccgggaact tccgagagcg ctacaccaga aagcggaccc 240

ggaaccagta ccgaacctag cgagggctct gctccgggca gcccagccgg ctctcctaca 300

tccacggagg agggcacttc cgaatccgcc accccggagt cagggccagg atctgaaccc 360

gctacctcag gcagtgagac gccaggaacg agcgagtccg ctacaccgga gagtgggcca 420

gggagccctg ctggatctcc tacgtccact gaggaagggt caccagcggg ctcgcccacc 480

agcactgaag aaggtgcctc gagcagtgat ctcggtgagc tgcctgtgga cgcaagattt 540

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gaaacaaaga actccttgat gcaggatagg gatgctgcat ctgctcgggc ctggcctaaa 1140

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tcagcacccg gctctgagcc ggccacaagt ggcagtgaga cacccggcac ttcagagagt 6480

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tcgagcgaac aaaaactcat ctcagaagag gatctgtcac ctgcaaccgg tcatcatcac 6600

catcaccatt ga 6612

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<211> 2203
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1285

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Gly Ser Ser
20 25 30

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
35 40 45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
50 55 60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65 70 75 80

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Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
                85                  90                  95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
            100                 105                 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
        115                 120                 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
    130                 135                 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145                 150                 155                 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350
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Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620
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Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895
```

```
Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155
```

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Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
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Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Gly Ala Pro Gly  Thr Pro Gly
    1310                1315                1320

Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1325                1330                1335

Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1340                1345                1350

Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly
    1355                1360                1365

Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly Pro Gly  Thr Pro Gly
    1370                1375                1380

Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1385                1390                1395

Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1400                1405                1410
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```
Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly
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Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr
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Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1445                 1450                 1455

Ser Thr  Gly Ser Pro Ala Ser  Ser Thr Phe Lys Glu  Asn Tyr Arg
    1460                 1465                 1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                 1480                 1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                 1495                 1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                 1510                 1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                 1525                 1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                 1540                 1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                 1555                 1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                 1570                 1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                 1585                 1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                 1600                 1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                 1615                 1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                 1630                 1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                 1645                 1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                 1660                 1665
```

```
Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920
```

```
Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                1960                1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                1975                1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                1990                1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                2005                2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                2020                2025

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    2030                2035                2040

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2045                2050                2055

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    2060                2065                2070

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    2075                2080                2085

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    2090                2095                2100

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    2105                2110                2115

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    2120                2125                2130

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Glu Pro Ala
    2135                2140                2145

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2150                2155                2160

Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
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Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Ala Thr Gly His His His His His His
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20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Gly Ala Pro Gly Ser Ser Pro Ser
35 40 45

Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
50 55 60

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
65 70 75 80

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Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro
                85                  90                  95

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
            100                 105                 110

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
        115                 120                 125

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser
    130                 135                 140

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
145                 150                 155                 160

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
                165                 170                 175

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350
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Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
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Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620
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Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895
```

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Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
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Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155
```

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Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
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Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Gly Ala Pro Gly  Thr Pro Gly
    1310                1315                1320

Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1325                1330                1335

Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1340                1345                1350

Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly
    1355                1360                1365

Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly Pro Gly  Thr Pro Gly
    1370                1375                1380

Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1385                1390                1395

Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1400                1405                1410
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Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly
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Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr
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Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1445                 1450                 1455

Ser Thr  Gly Ser Pro Ala Ser  Ser Thr Phe Lys Glu  Asn Tyr Arg
    1460                 1465                 1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                 1480                 1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                 1495                 1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                 1510                 1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                 1525                 1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                 1540                 1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                 1555                 1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                 1570                 1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                 1585                 1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                 1600                 1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                 1615                 1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                 1630                 1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                 1645                 1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                 1660                 1665
```

```
Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920
```

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Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                1960                1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                1975                1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                1990                1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                2005                2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                2020                2025

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    2030                2035                2040

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2045                2050                2055

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    2060                2065                2070

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    2075                2080                2085

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    2090                2095                2100

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    2105                2110                2115

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    2120                2125                2130

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Glu Pro Ala
    2135                2140                2145

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2150                2155                2160

Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
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Pro Ala Thr Gly His His His His His His
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35 40 45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
50 55 60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65 70 75 80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser

85 90 95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
100 105 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
115 120 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
130 135 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145 150 155 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
260 265 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
275 280 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
290 295 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305 310 315 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
325 330 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
340 345 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln

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385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
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625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
645 650 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
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Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
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Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
885 890 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser

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Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
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Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
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Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
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Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
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Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
995 1000 1005

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Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
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Ser Ala Pro Gly Ala Ser Ser Pro Pro Val Leu Lys Arg His Gln
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Ala Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
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Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
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Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln
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Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp
1100 1105 1110

Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala
1115 1120 1125

Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu
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Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
1145 1150 1155

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val

1160 1165 1170

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1175 1180 1185

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1190 1195 1200

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
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Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
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Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1235 1240 1245

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1250 1255 1260

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1265 1270 1275

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1280 1285 1290

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1295 1300 1305

Cys Asn Ile Gln Met Glu Asp Pro Gly Ala Pro Gly Thr Pro Gly
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Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser
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Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
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Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
1370 1375 1380

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser
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Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
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Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly

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Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
1430 1435 1440

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
1445 1450 1455

Ser Thr Gly Ser Pro Ala Ser Ser Thr Phe Lys Glu Asn Tyr Arg
1460 1465 1470

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1475 1480 1485

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1490 1495 1500

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
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Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1520 1525 1530

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1535 1540 1545

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1550 1555 1560

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1565 1570 1575

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1580 1585 1590

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1595 1600 1605

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1610 1615 1620

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1625 1630 1635

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1640 1645 1650

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1655 1660 1665

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe

1670 1675 1680

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1685 1690 1695

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1700 1705 1710

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1715 1720 1725

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1730 1735 1740

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1745 1750 1755

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1760 1765 1770

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1775 1780 1785

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1790 1795 1800

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1805 1810 1815

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1820 1825 1830

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1835 1840 1845

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1850 1855 1860

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1865 1870 1875

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Pro Gly Thr
1880 1885 1890

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1895 1900 1905

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
1910 1915 1920

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr

1925 1930 1935

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1940 1945 1950

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
1955 1960 1965

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
1970 1975 1980

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1985 1990 1995

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2000 2005 2010

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
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Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
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Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
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Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
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Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
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Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
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Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
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Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
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Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
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Pro Ala Thr Gly His His His His His His
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

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Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365
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Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gly Ala Pro Thr Ser Thr Glu
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Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
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Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
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Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
    450                 455                 460

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
465                 470                 475                 480

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
                485                 490                 495

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
            500                 505                 510

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
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Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
    530                 535                 540

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640
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Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910
```

```
Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
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Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
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Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
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Gln Gly  Ala Pro Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
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Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser
    1145                1150                1155

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
    1160                1165                1170
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Ser Gly  Ser Glu Thr Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
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Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
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Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Pro
    1205                1210                1215

Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Ser Pro  Ala Gly Ser
    1220                1225                1230

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1235                1240                1245

Thr Glu  Glu Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1250                1255                1260

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Ala Ser
    1265                1270                1275

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1280                1285                1290

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1295                1300                1305

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1310                1315                1320

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1325                1330                1335

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1340                1345                1350

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1355                1360                1365

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1370                1375                1380

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1385                1390                1395

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1400                1405                1410

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1415                1420                1425
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Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1430                1435                1440

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1445                1450                1455

Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1460                1465                1470

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1475                1480                1485

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1490                1495                1500

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1505                1510                1515

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1520                1525                1530

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1535                1540                1545

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1550                1555                1560

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1565                1570                1575

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1580                1585                1590

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1595                1600                1605

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1610                1615                1620

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1625                1630                1635

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1640                1645                1650

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1655                1660                1665

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1670                1675                1680
```

```
Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1685                1690                1695

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1700                1705                1710

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1715                1720                1725

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1730                1735                1740

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1745                1750                1755

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1760                1765                1770

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1775                1780                1785

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1790                1795                1800

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1805                1810                1815

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1820                1825                1830

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1835                1840                1845

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    1850                1855                1860

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    1865                1870                1875

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    1880                1885                1890

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1895                1900                1905

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    1910                1915                1920

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1925                1930                1935
```

```
Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1940                1945                1950

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    1955                1960                1965

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    1970                1975                1980

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1985                1990                1995

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2000                2005                2010

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser
    2015                2020                2025

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Thr Glu
    2030                2035                2040

Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2045                2050                2055

Glu Ser  Gly Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2060                2065                2070

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2075                2080                2085

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Ser  Glu Pro Ala
    2090                2095                2100

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Pro Ala Gly  Ser Pro Thr
    2105                2110                2115

Ser Thr  Glu Glu Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    2120                2125                2130

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    2135                2140                2145

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    2150                2155                2160

Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu
    2165                2170                2175

Gly Ser  Ala Pro Ala Ser Ser  Glu Gln Lys Leu Ile  Ser Glu Glu
    2180                2185                2190
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

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Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365
```

```
Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
                405                 410                 415

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser
            420                 425                 430

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly
        435                 440                 445

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
    450                 455                 460

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala
465                 470                 475                 480

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly
                485                 490                 495

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
            500                 505                 510

Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala
        515                 520                 525

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
    530                 535                 540

Ser Ser Thr Gly Ser Pro Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640
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Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910
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Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
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Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
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Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
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Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
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Gln Gly  Ala Pro Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
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Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser
    1145                1150                1155

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
    1160                1165                1170
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Ser Gly  Ser Glu Thr Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
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Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
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Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Pro
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Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Ser Pro  Ala Gly Ser
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Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
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Thr Glu  Glu Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
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Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Ala Ser
    1265                1270                1275

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1280                1285                1290

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1295                1300                1305

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
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Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
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Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
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Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
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Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
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Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
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Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
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Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1415                1420                1425
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Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
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Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1445                1450                1455

Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1460                1465                1470

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1475                1480                1485

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1490                1495                1500

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1505                1510                1515

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1520                1525                1530

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1535                1540                1545

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1550                1555                1560

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1565                1570                1575

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1580                1585                1590

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1595                1600                1605

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1610                1615                1620

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
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Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1640                1645                1650

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1655                1660                1665

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
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Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
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Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1700                 1705                 1710

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1715                 1720                 1725

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1730                 1735                 1740

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1745                 1750                 1755

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1760                 1765                 1770

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
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Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1790                 1795                 1800

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1805                 1810                 1815

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1820                 1825                 1830

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1835                 1840                 1845

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    1850                 1855                 1860

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    1865                 1870                 1875

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    1880                 1885                 1890

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1895                 1900                 1905

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
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Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1925                 1930                 1935
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Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
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Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    1955                1960                1965

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    1970                1975                1980

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1985                1990                1995

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2000                2005                2010

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser
    2015                2020                2025

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Thr Glu
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Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2045                2050                2055

Glu Ser  Gly Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2060                2065                2070

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2075                2080                2085

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Ser  Glu Pro Ala
    2090                2095                2100

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Pro Ala Gly  Ser Pro Thr
    2105                2110                2115

Ser Thr  Glu Glu Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    2120                2125                2130

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    2135                2140                2145

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    2150                2155                2160

Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu
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Gly Ser  Ala Pro Ala Ser Ser  Glu Gln Lys Leu Ile  Ser Glu Glu
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

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Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365
```

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Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gly Ala Pro Thr Ser Thr Glu
                405                 410                 415

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
            420                 425                 430

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
        435                 440                 445

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
    450                 455                 460

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
465                 470                 475                 480

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
                485                 490                 495

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
            500                 505                 510

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
        515                 520                 525

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
    530                 535                 540

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640
```

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Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910
```

```
Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
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Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Gly  Ala Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1130                1135                1140

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1145                1150                1155

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1160                1165                1170
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```
Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
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Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
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Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1205                1210                1215

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1220                1225                1230

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1235                1240                1245

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1250                1255                1260

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
    1265                1270                1275

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1280                1285                1290

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1295                1300                1305

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1310                1315                1320

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1325                1330                1335

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1340                1345                1350

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1355                1360                1365

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1370                1375                1380

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1385                1390                1395

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1400                1405                1410

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1415                1420                1425
```

```
Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
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Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1445                1450                1455

Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1460                1465                1470

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1475                1480                1485

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1490                1495                1500

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1505                1510                1515

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1520                1525                1530

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1535                1540                1545

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1550                1555                1560

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1565                1570                1575

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1580                1585                1590

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1595                1600                1605

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1610                1615                1620

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1625                1630                1635

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1640                1645                1650

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1655                1660                1665

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1670                1675                1680
```

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Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
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Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1700                 1705                 1710

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1715                 1720                 1725

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1730                 1735                 1740

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1745                 1750                 1755

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1760                 1765                 1770

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1775                 1780                 1785

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1790                 1795                 1800

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1805                 1810                 1815

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1820                 1825                 1830

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1835                 1840                 1845

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    1850                 1855                 1860

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    1865                 1870                 1875

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    1880                 1885                 1890

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1895                 1900                 1905

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
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Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1925                 1930                 1935
```

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Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
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Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    1955                1960                1965

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    1970                1975                1980

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1985                1990                1995

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2000                2005                2010

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser
    2015                2020                2025

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Thr Glu
    2030                2035                2040

Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2045                2050                2055

Glu Ser  Gly Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2060                2065                2070

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2075                2080                2085

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Ser  Glu Pro Ala
    2090                2095                2100

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Pro Ala Gly  Ser Pro Thr
    2105                2110                2115

Ser Thr  Glu Glu Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    2120                2125                2130

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    2135                2140                2145

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    2150                2155                2160

Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu
    2165                2170                2175

Gly Ser  Ala Pro Ala Ser Ser  Glu Gln Lys Leu Ile  Ser Glu Glu
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Asp Leu Ser Pro Ala Thr Gly His His His His His His
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<223> Description of Artificial Sequence: Synthetic polynucleotide

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tcagagagtg ccacccccga gagtggccca ggcactagta ccgagccctc tgaaggcagt 6540

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20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala

100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His

370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
405 410 415

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
420 425 430

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
435 440 445

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
450 455 460

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
465 470 475 480

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
485 490 495

Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
500 505 510

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
515 520 525

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
530 535 540

Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545 550 555 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
565 570 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
580 585 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
595 600 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
610 615 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly

645 650 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
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Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
835 840 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
885 890 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
900 905 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser

915 920 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
930 935 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945 950 955 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
965 970 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
980 985 990

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
995 1000 1005

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser
1010 1015 1020

Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
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Ser Ala Pro Gly Ala Ser Ser Pro Pro Val Leu Lys Arg His Gln
1040 1045 1050

Ala Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
1055 1060 1065

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
1070 1075 1080

Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln
1085 1090 1095

Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp
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Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala
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Gln Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
1130 1135 1140

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
1145 1150 1155

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
1160 1165 1170

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr

1175 1180 1185

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
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Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
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Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
1220 1225 1230

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
1235 1240 1245

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
1250 1255 1260

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser
1265 1270 1275

Ser Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val
1280 1285 1290

Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg
1295 1300 1305

Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
1310 1315 1320

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala
1325 1330 1335

Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu
1340 1345 1350

Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro
1355 1360 1365

Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala
1370 1375 1380

Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser
1385 1390 1395

Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
1400 1405 1410

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg
1415 1420 1425

Gln Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp

1430 1435 1440

Glu Thr Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys
1445 1450 1455

Arg Ala Pro Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu
1460 1465 1470

Asn Tyr Arg Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu
1475 1480 1485

Pro Gly Leu Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu
1490 1495 1500

Leu Ser Met Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser
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Gly His Val Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala
1520 1525 1530

Leu Tyr Asn Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu
1535 1540 1545

Pro Ser Lys Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu
1550 1555 1560

His Leu His Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn
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Lys Cys Gln Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp
1580 1585 1590

Phe Gln Ile Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys
1595 1600 1605

Leu Ala Arg Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr
1610 1615 1620

Lys Glu Pro Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met
1625 1630 1635

Ile Ile His Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser
1640 1645 1650

Ser Leu Tyr Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly
1655 1660 1665

Lys Lys Trp Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met
1670 1675 1680

Val Phe Phe Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile

1685 1690 1695

Phe Asn Pro Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr
1700 1705 1710

His Tyr Ser Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys
1715 1720 1725

Asp Leu Asn Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala
1730 1735 1740

Ile Ser Asp Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met
1745 1750 1755

Phe Ala Thr Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly
1760 1765 1770

Arg Ser Asn Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp
1775 1780 1785

Leu Gln Val Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr
1790 1795 1800

Thr Gln Gly Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu
1805 1810 1815

Phe Leu Ile Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe
1820 1825 1830

Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser
1835 1840 1845

Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg
1850 1855 1860

Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu
1865 1870 1875

Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala
1880 1885 1890

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
1895 1900 1905

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
1910 1915 1920

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
1925 1930 1935

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly

1940 1945 1950

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
1955 1960 1965

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
1970 1975 1980

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
1985 1990 1995

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2000 2005 2010

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser
2015 2020 2025

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu
2030 2035 2040

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
2045 2050 2055

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2060 2065 2070

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2075 2080 2085

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala
2090 2095 2100

Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
2105 2110 2115

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2120 2125 2130

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
2135 2140 2145

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
2150 2155 2160

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
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Gly Ser Ala Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu
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atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

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tcagagagtg ccacccccga gagtggccca ggcactagta ccgagccctc tgaaggcagt 6540

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20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

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Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380
```

```
Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
                405                 410                 415

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser
            420                 425                 430

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly
        435                 440                 445

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
    450                 455                 460

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala
465                 470                 475                 480

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly
                485                 490                 495

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
            500                 505                 510

Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala
        515                 520                 525

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
    530                 535                 540

Ser Ser Thr Gly Ser Pro Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655
```

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Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925
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Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Gly  Ala Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1130                1135                1140

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1145                1150                1155

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1160                1165                1170

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1175                1180                1185
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Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
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Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1205                1210                1215

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1220                1225                1230

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1235                1240                1245

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1250                1255                1260

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
    1265                1270                1275

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1280                1285                1290

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1295                1300                1305

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1310                1315                1320

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1325                1330                1335

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1340                1345                1350

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1355                1360                1365

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1370                1375                1380

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1385                1390                1395

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1400                1405                1410

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1415                1420                1425

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1430                1435                1440
```

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Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
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Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1460                1465                1470

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1475                1480                1485

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1490                1495                1500

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1505                1510                1515

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1520                1525                1530

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1535                1540                1545

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1550                1555                1560

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1565                1570                1575

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1580                1585                1590

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1595                1600                1605

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1610                1615                1620

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1625                1630                1635

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1640                1645                1650

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1655                1660                1665

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1670                1675                1680

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1685                1690                1695
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Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
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His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1715                 1720                 1725

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1730                 1735                 1740

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1745                 1750                 1755

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1760                 1765                 1770

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1775                 1780                 1785

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1790                 1795                 1800

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1805                 1810                 1815

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1820                 1825                 1830

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1835                 1840                 1845

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    1850                 1855                 1860

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    1865                 1870                 1875

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    1880                 1885                 1890

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1895                 1900                 1905

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    1910                 1915                 1920

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1925                 1930                 1935

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1940                 1945                 1950
```

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Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    1955                 1960                 1965

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    1970                 1975                 1980

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1985                 1990                 1995

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2000                 2005                 2010

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser
    2015                 2020                 2025

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Thr Glu
    2030                 2035                 2040

Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2045                 2050                 2055

Glu Ser  Gly Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2060                 2065                 2070

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2075                 2080                 2085

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Ser  Glu Pro Ala
    2090                 2095                 2100

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Pro Ala Gly  Ser Pro Thr
    2105                 2110                 2115

Ser Thr  Glu Glu Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    2120                 2125                 2130

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    2135                 2140                 2145

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    2150                 2155                 2160

Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu
    2165                 2170                 2175

Gly Ser  Ala Pro Ala Ser Ser  Glu Gln Lys Leu Ile  Ser Glu Glu
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Asp Leu  Ser Pro Ala Thr Gly  His His His His His  His
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

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Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
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Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380
```

```
Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
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Ala Pro Asp Gly Ala Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
                405                 410                 415

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser
            420                 425                 430

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly
        435                 440                 445

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
    450                 455                 460

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala
465                 470                 475                 480

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly
                485                 490                 495

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
            500                 505                 510

Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala
        515                 520                 525

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
    530                 535                 540

Ser Ser Thr Gly Ser Pro Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655
```

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Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
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Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925
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Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
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Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
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Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
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```

```
Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
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Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
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Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                 1225                 1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                 1240                 1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                 1255                 1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                 1270                 1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                 1285                 1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1295                 1300                 1305

Pro Ser  Pro Ala Gly Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser
    1310                 1315                 1320

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro
    1325                 1330                 1335

Ser Glu  Gly Ser Ala Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1340                 1345                 1350

Thr Glu  Glu Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1355                 1360                 1365

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1370                 1375                 1380

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
    1385                 1390                 1395

Ser Gly  Ser Glu Thr Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1400                 1405                 1410

Glu Thr  Pro Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1415                 1420                 1425

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1430                 1435                 1440
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Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ala Ser  Ser Ala Pro
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Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                1555                1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                1615                1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695
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Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
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Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950
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Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
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Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                 1975                 1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                 1990                 1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                 2005                 2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                 2020                 2025

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    2030                 2035                 2040

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2045                 2050                 2055

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    2075                 2080                 2085

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    2090                 2095                 2100

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    2105                 2110                 2115

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    2120                 2125                 2130

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Glu Pro Ala
    2135                 2140                 2145

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2150                 2155                 2160

Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    2165                 2170                 2175

Pro Ala  Ser Ser Glu Gln Lys  Leu Ile Ser Glu Glu  Asp Leu Ser
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Pro Ala  Thr Gly His His His  His His His
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

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Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380
```

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Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
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Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
                405                 410                 415

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            420                 425                 430

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
        435                 440                 445

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    450                 455                 460

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
465                 470                 475                 480

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
                485                 490                 495

Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            500                 505                 510

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
        515                 520                 525

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
    530                 535                 540

Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655
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Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925
```

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Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
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Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185
```

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Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
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Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
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Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                 1225                 1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                 1240                 1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                 1255                 1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                 1270                 1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                 1285                 1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1295                 1300                 1305

Pro Ser  Pro Ala Gly Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser
    1310                 1315                 1320

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro
    1325                 1330                 1335

Ser Glu  Gly Ser Ala Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1340                 1345                 1350

Thr Glu  Glu Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1355                 1360                 1365

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1370                 1375                 1380

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
    1385                 1390                 1395

Ser Gly  Ser Glu Thr Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1400                 1405                 1410

Glu Thr  Pro Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1415                 1420                 1425

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
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Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ala Ser  Ser Ala Pro
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Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                1555                1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                1615                1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695
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Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
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Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950
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Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                 1960                 1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                 1975                 1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                 1990                 1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                 2005                 2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                 2020                 2025

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    2030                 2035                 2040

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2045                 2050                 2055

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    2060                 2065                 2070

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    2075                 2080                 2085

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    2090                 2095                 2100

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    2105                 2110                 2115

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    2120                 2125                 2130

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Glu Pro Ala
    2135                 2140                 2145

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2150                 2155                 2160

Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    2165                 2170                 2175

Pro Ala  Ser Ser Glu Gln Lys  Leu Ile Ser Glu Glu  Asp Leu Ser
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Pro Ala  Thr Gly His His His  His His His
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
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Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val

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Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
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Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
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His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
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Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
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Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
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Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
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Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
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Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
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Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
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Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
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Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
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Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gly Ala Pro Thr Ser Thr Glu
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Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
435 440 445

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Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
465 470 475 480

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
485 490 495

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
500 505 510

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
515 520 525

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
530 535 540

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Gln
545 550 555 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
565 570 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
580 585 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
595 600 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
610 615 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
645 650 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr

660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
835 840 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
885 890 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
900 905 910

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915 920 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly

930 935 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945 950 955 960

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Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
980 985 990

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
995 1000 1005

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Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
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Ala Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
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Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
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Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln
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Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp
1100 1105 1110

Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala
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Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu
1130 1135 1140

Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
1145 1150 1155

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1160 1165 1170

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1175 1180 1185

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg

1190 1195 1200

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
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Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
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Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1235 1240 1245

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1250 1255 1260

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1265 1270 1275

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1280 1285 1290

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Gly Ala
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Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro
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1355 1360 1365

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
1370 1375 1380

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr
1385 1390 1395

Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
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Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
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Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Ala Pro

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Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
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Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
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Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
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Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
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Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
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Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
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Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
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Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
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Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
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Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                1615                1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
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Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
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Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
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Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
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Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
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Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1730 1735 1740

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1745 1750 1755

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1760 1765 1770

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1775 1780 1785

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
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Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
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Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1835 1840 1845

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
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1865 1870 1875

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Pro Gly Thr
1880 1885 1890

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1895 1900 1905

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
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Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly

1955 1960 1965

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
1970 1975 1980

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1985 1990 1995

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2000 2005 2010

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
2015 2020 2025

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Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

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Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
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Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400
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Ala Pro Asp Gly Ala Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
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Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser
            420                 425                 430

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly
        435                 440                 445

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
    450                 455                 460

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala
465                 470                 475                 480

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly
                485                 490                 495

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
            500                 505                 510

Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala
        515                 520                 525

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
    530                 535                 540

Ser Ser Thr Gly Ser Pro Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670
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Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940
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Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
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Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200
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Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
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Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                 1225                 1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                 1240                 1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                 1255                 1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                 1270                 1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                 1285                 1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1295                 1300                 1305

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala
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Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1325                 1330                 1335

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
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Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
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Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala
    1370                 1375                 1380

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1385                 1390                 1395

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1400                 1405                 1410

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
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Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ala
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Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Ala Ser  Ser Ala Pro
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Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
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Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                 1480                 1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                 1495                 1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                 1510                 1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                 1525                 1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                 1540                 1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                 1555                 1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                 1570                 1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                 1585                 1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                 1600                 1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                 1615                 1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                 1630                 1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                 1645                 1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                 1660                 1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                 1675                 1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                 1690                 1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                 1705                 1710
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Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
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Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
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Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
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Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                1960                1965
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Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                1975                1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                1990                1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                2005                2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                2020                2025

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    2030                2035                2040

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    2075                2080                2085

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    2090                2095                2100

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    2105                2110                2115

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    2120                2125                2130

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Glu Pro Ala
    2135                2140                2145

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2150                2155                2160

Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    2165                2170                2175

Pro Ala  Ser Ser Glu Gln Lys  Leu Ile Ser Glu Glu  Asp Leu Ser
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Pro Ala  Thr Gly His His His  His His His
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

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Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
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Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400
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Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            420                 425                 430

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
        435                 440                 445

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    450                 455                 460

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
465                 470                 475                 480

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
                485                 490                 495

Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            500                 505                 510

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
        515                 520                 525

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
    530                 535                 540

Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670
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Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940
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Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
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Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
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Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
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Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                 1225                 1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                 1240                 1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                 1255                 1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                 1270                 1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                 1285                 1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1295                 1300                 1305

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala
    1310                 1315                 1320

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1325                 1330                 1335

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
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Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
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Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala
    1370                 1375                 1380

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1385                 1390                 1395

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1400                 1405                 1410

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
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Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ala
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Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Ala Ser  Ser Ala Pro
    1445                 1450                 1455
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Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
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Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                 1480                 1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                 1495                 1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                 1510                 1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                 1525                 1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                 1540                 1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                 1555                 1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                 1570                 1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                 1585                 1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                 1600                 1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                 1615                 1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                 1630                 1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                 1645                 1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                 1660                 1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                 1675                 1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                 1690                 1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                 1705                 1710
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Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
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Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                1960                1965
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Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
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Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                2005                2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                2020                2025

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    2030                2035                2040

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    2075                2080                2085

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    2090                2095                2100

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
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Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    2120                2125                2130

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Glu Pro Ala
    2135                2140                2145

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2150                2155                2160

Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    2165                2170                2175

Pro Ala  Ser Ser Glu Gln Lys  Leu Ile Ser Glu Glu  Asp Leu Ser
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

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Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400
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Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gly Ala Pro Thr Ser Thr Glu
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Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
           420                 425                 430

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
       435                 440                 445

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
   450                 455                 460

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
465                 470                 475                 480

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
               485                 490                 495

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
           500                 505                 510

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
       515                 520                 525

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
   530                 535                 540

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
               565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
           580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
       595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
   610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
               645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
           660                 665                 670
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Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940
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Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
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Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
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Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
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Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                 1225                 1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                 1240                 1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                 1255                 1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                 1270                 1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                 1285                 1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1295                 1300                 1305

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala
    1310                 1315                 1320

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1325                 1330                 1335

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
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Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
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Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala
    1370                 1375                 1380

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1385                 1390                 1395

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1400                 1405                 1410

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
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Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ala
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Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Ala Ser  Ser Ala Pro
    1445                 1450                 1455
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Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
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Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                 1480                 1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                 1495                 1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                 1510                 1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                 1525                 1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                 1540                 1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                 1555                 1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                 1570                 1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                 1585                 1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                 1600                 1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                 1615                 1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                 1630                 1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                 1645                 1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                 1660                 1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                 1675                 1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                 1690                 1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                 1705                 1710
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Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
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Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
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Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
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Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
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Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
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Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
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Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
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Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
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Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
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Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                1960                1965
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Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
2015 2020 2025

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn

130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gly Ala Pro Thr Ser Thr Glu

405 410 415

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
420 425 430

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
435 440 445

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
450 455 460

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
465 470 475 480

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
485 490 495

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
500 505 510

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
515 520 525

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
530 535 540

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Gln
545 550 555 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
565 570 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
580 585 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
595 600 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
610 615 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
645 650 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn

675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
835 840 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
885 890 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
900 905 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
915 920 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
930 935 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser

945 950 955 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
965 970 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
980 985 990

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
995 1000 1005

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser
1010 1015 1020

Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
1025 1030 1035

Ser Ala Pro Gly Ala Ser Ser Pro Pro Val Leu Lys Arg His Gln
1040 1045 1050

Ala Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
1055 1060 1065

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
1070 1075 1080

Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln
1085 1090 1095

Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp
1100 1105 1110

Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala
1115 1120 1125

Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu
1130 1135 1140

Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
1145 1150 1155

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1160 1165 1170

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1175 1180 1185

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1190 1195 1200

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr

1205 1210 1215

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1220 1225 1230

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1235 1240 1245

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1250 1255 1260

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1265 1270 1275

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1280 1285 1290

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1295 1300 1305

Cys Asn Ile Gly Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr
1310 1315 1320

Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
1325 1330 1335

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Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
1355 1360 1365

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
1370 1375 1380

Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
1385 1390 1395

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
1400 1405 1410

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro
1415 1420 1425

Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
1430 1435 1440

Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
1445 1450 1455

Ala Ser Ser Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg

1460 1465 1470

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1475 1480 1485

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1490 1495 1500

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1505 1510 1515

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1520 1525 1530

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1535 1540 1545

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1550 1555 1560

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1565 1570 1575

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1580 1585 1590

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1595 1600 1605

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1610 1615 1620

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
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Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1640 1645 1650

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1655 1660 1665

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1670 1675 1680

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1685 1690 1695

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
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Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn

1715 1720 1725

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1730 1735 1740

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1745 1750 1755

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1760 1765 1770

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
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Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1790 1795 1800

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1805 1810 1815

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1820 1825 1830

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1835 1840 1845

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1850 1855 1860

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1865 1870 1875

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Pro Gly Thr
1880 1885 1890

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1895 1900 1905

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
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1925 1930 1935

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1940 1945 1950

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
1955 1960 1965

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro

1970 1975 1980

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1985 1990 1995

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2000 2005 2010

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
2015 2020 2025

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
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Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2045 2050 2055

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
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2075 2080 2085

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
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Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
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Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
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catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

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tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

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Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
                405                 410                 415
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Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser
            420                 425                 430

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly
        435                 440                 445

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
    450                 455                 460

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala
465                 470                 475                 480

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly
                485                 490                 495

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
            500                 505                 510

Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala
        515                 520                 525

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
    530                 535                 540

Ser Ser Thr Gly Ser Pro Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685
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Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960
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Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
               965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
           980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215
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Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
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Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                 1240                 1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                 1255                 1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                 1270                 1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                 1285                 1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                 1300                 1305

Cys Asn  Ile Gly Ala Pro Ser  Pro Ala Gly Ser Pro  Thr Ser Thr
    1310                 1315                 1320

Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    1325                 1330                 1335

Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala Pro Gly  Ser Pro Ala
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Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr Ser Thr  Glu Pro Ser
    1355                 1360                 1365

Glu Gly  Ser Ala Pro Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    1370                 1375                 1380

Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    1385                 1390                 1395

Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr Pro Gly  Ser Glu Pro
    1400                 1405                 1410

Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Ser Pro Ala  Gly Ser Pro
    1415                 1420                 1425

Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1430                 1435                 1440

Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    1445                 1450                 1455

Ala Ser  Ser Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                 1465                 1470
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Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
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Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                 1495                 1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                 1510                 1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                 1525                 1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                 1540                 1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                 1555                 1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                 1570                 1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                 1585                 1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                 1600                 1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                 1615                 1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                 1630                 1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                 1645                 1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                 1660                 1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                 1675                 1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                 1690                 1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
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Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
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Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
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Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
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Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
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Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
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Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
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Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
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Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                1960                1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                1975                1980
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Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                 2005                 2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                 2020                 2025

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
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Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    2075                 2080                 2085

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
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Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
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Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    2120                 2125                 2130

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Glu Pro Ala
    2135                 2140                 2145

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2150                 2155                 2160

Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    2165                 2170                 2175

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Pro Ala  Thr Gly His His His  His His His
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

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Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
                405                 410                 415
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Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            420                 425                 430

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
        435                 440                 445

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    450                 455                 460

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
465                 470                 475                 480

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
                485                 490                 495

Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            500                 505                 510

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
        515                 520                 525

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
    530                 535                 540

Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685
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Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960
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Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
               965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
           980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
       995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215
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Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                 1225                 1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                 1240                 1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                 1255                 1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                 1270                 1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                 1285                 1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                 1300                 1305

Cys Asn  Ile Gly Ala Pro Ser  Pro Ala Gly Ser Pro  Thr Ser Thr
    1310                 1315                 1320

Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    1325                 1330                 1335

Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala Pro Gly  Ser Pro Ala
    1340                 1345                 1350

Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr Ser Thr  Glu Pro Ser
    1355                 1360                 1365

Glu Gly  Ser Ala Pro Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    1370                 1375                 1380

Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    1385                 1390                 1395

Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr Pro Gly  Ser Glu Pro
    1400                 1405                 1410

Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Ser Pro Ala  Gly Ser Pro
    1415                 1420                 1425

Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1430                 1435                 1440

Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    1445                 1450                 1455

Ala Ser  Ser Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                 1465                 1470
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Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                1555                1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                1615                1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725
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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
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Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                1960                1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                1975                1980
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1985 1990 1995

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2000 2005 2010

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
2015 2020 2025

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
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Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2045 2050 2055

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
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Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
2075 2080 2085

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
2090 2095 2100

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
2105 2110 2115

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
2120 2125 2130

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
2135 2140 2145

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2150 2155 2160

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2165 2170 2175

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
2180 2185 2190

Pro Ala Thr Gly His His His His His His
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<211> 6612
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polynucleotide

<400> 1318

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<211> 2203
<212> PRT
<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1319

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

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Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
                405                 410                 415
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Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser
            420                 425                 430

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly
        435                 440                 445

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
    450                 455                 460

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala
465                 470                 475                 480

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly
                485                 490                 495

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
            500                 505                 510

Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala
        515                 520                 525

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
    530                 535                 540

Ser Ser Thr Gly Ser Pro Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685
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Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960
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Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
               965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
           980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
       995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215
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Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gly Ala Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    1310                1315                1320

Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1325                1330                1335

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1340                1345                1350

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala
    1355                1360                1365

Ser Thr  Gly Thr Gly Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    1370                1375                1380

Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1385                1390                1395

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1400                1405                1410

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser
    1415                1420                1425

Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    1430                1435                1440

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1445                1450                1455

Ala Ser  Ser Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470
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Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                1555                1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                1615                1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725
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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                1960                1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                1975                1980
```

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1985 1990 1995

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2000 2005 2010

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
2015 2020 2025

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
2030 2035 2040

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2045 2050 2055

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
2060 2065 2070

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
2075 2080 2085

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
2090 2095 2100

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
2105 2110 2115

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
2120 2125 2130

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
2135 2140 2145

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2150 2155 2160

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2165 2170 2175

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
2180 2185 2190

Pro Ala Thr Gly His His His His His His
2195 2200

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atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

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gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

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tcagcacccg gctctgagcc ggccacaagt ggcagtgaga cacccggcac ttcagagagt 6480

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catcaccatt ga 6612

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<211> 2203
<212> PRT
<213> Artificial Sequence

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<400> 1321

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser

145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gly Ala Pro Thr Ser Thr Glu
405 410 415

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser

420 425 430

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
435 440 445

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
450 455 460

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
465 470 475 480

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
485 490 495

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
500 505 510

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
515 520 525

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
530 535 540

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Gln
545 550 555 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
565 570 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
580 585 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
595 600 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
610 615 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
645 650 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys

690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
755 760 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
770 775 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785 790 795 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
805 810 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
820 825 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
835 840 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
850 855 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865 870 875 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
885 890 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
900 905 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
915 920 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
930 935 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945 950 955 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser

965 970 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
980 985 990

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
995 1000 1005

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser
1010 1015 1020

Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
1025 1030 1035

Ser Ala Pro Gly Ala Ser Ser Pro Pro Val Leu Lys Arg His Gln
1040 1045 1050

Ala Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
1055 1060 1065

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
1070 1075 1080

Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln
1085 1090 1095

Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp
1100 1105 1110

Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala
1115 1120 1125

Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu
1130 1135 1140

Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
1145 1150 1155

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1160 1165 1170

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1175 1180 1185

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1190 1195 1200

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1205 1210 1215

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys

1220 1225 1230

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1235 1240 1245

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1250 1255 1260

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1265 1270 1275

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1280 1285 1290

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
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Cys Asn Ile Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
1310 1315 1320

Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
1325 1330 1335

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
1340 1345 1350

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala
1355 1360 1365

Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
1370 1375 1380

Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
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Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
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Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser
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Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
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Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
1445 1450 1455

Ala Ser Ser Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1460 1465 1470

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu

1475 1480 1485

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1490 1495 1500

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1505 1510 1515

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1520 1525 1530

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1535 1540 1545

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1550 1555 1560

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1565 1570 1575

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1580 1585 1590

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1595 1600 1605

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1610 1615 1620

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1625 1630 1635

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1640 1645 1650

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1655 1660 1665

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1670 1675 1680

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1685 1690 1695

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1700 1705 1710

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1715 1720 1725

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp

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Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                1960                1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                1975                1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
```

1985 1990 1995

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2000 2005 2010

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
2015 2020 2025

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
2030 2035 2040

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2045 2050 2055

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
2060 2065 2070

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
2075 2080 2085

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
2090 2095 2100

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
2105 2110 2115

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
2120 2125 2130

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
2135 2140 2145

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2150 2155 2160

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Ala Thr Gly His His His His His His
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<211> 6612
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tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

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<211> 2203
<212> PRT
<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

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His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
                405                 410                 415

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            420                 425                 430
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Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
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Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    450                 455                 460

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
465                 470                 475                 480

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
                485                 490                 495

Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            500                 505                 510

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
        515                 520                 525

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
    530                 535                 540

Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700
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Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975
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Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
           980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
       995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230
```

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Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
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Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gly Ala Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    1310                1315                1320

Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1325                1330                1335

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1340                1345                1350

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala
    1355                1360                1365

Ser Thr  Gly Thr Gly Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
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Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1385                1390                1395

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1400                1405                1410

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser
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Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    1430                1435                1440

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1445                1450                1455

Ala Ser  Ser Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485
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Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
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Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                 1510                 1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                 1525                 1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                 1540                 1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                 1555                 1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                 1570                 1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                 1585                 1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                 1600                 1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                 1615                 1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                 1630                 1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                 1645                 1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                 1660                 1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                 1675                 1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                 1690                 1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                 1705                 1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                 1720                 1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                 1735                 1740
```

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Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                1960                1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                1975                1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                1990                1995
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Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2000 2005 2010

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
2015 2020 2025

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
2030 2035 2040

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2045 2050 2055

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
2060 2065 2070

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
2075 2080 2085

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
2090 2095 2100

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
2105 2110 2115

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
2120 2125 2130

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
2135 2140 2145

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2150 2155 2160

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2165 2170 2175

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Ala Thr Gly His His His His His His
2195 2200

<210> 1324
<211> 6612
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 1324

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<210> 1325
<211> 2203
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1325

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

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His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gly Ala Pro Thr Ser Thr Glu
                405                 410                 415

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
            420                 425                 430
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Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
        435                 440                 445

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
    450                 455                 460

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
465                 470                 475                 480

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
                485                 490                 495

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
            500                 505                 510

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
        515                 520                 525

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
    530                 535                 540

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700
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Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975
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Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230
```

```
Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Gly Ala Pro Ser  Pro Ala Gly
    1310                1315                1320

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1325                1330                1335

Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    1340                1345                1350

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr
    1355                1360                1365

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Thr  Ser Thr Glu
    1370                1375                1380

Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1385                1390                1395

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1400                1405                1410

Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Ser
    1415                1420                1425

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    1430                1435                1440

Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu
    1445                1450                1455

Gly Ser  Ala Pro Gly Ala Ser  Ser Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485
```

```
Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                 1495                 1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                 1510                 1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                 1525                 1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                 1540                 1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                 1555                 1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                 1570                 1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                 1585                 1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                 1600                 1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                 1615                 1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                 1630                 1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                 1645                 1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                 1660                 1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                 1675                 1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                 1690                 1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                 1705                 1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                 1720                 1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                 1735                 1740
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Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                1960                1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                1975                1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                1990                1995
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Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2000 2005 2010

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
2015 2020 2025

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
2030 2035 2040

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2045 2050 2055

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
2060 2065 2070

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
2075 2080 2085

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
2090 2095 2100

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
2105 2110 2115

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
2120 2125 2130

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
2135 2140 2145

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2150 2155 2160

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2165 2170 2175

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Ala Thr Gly His His His His His His
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<211> 6612
<212> DNA
<213> Artificial Sequence

<220>
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catcaccatt ga 6612

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<211> 2203
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

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His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
                405                 410                 415

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            420                 425                 430
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Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
        435                 440                 445

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    450                 455                 460

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
465                 470                 475                 480

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
                485                 490                 495

Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            500                 505                 510

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
        515                 520                 525

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
    530                 535                 540

Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700
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Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975
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Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230
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Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
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Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Gly Ala Pro Ser  Pro Ala Gly
    1310                1315                1320

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1325                1330                1335

Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    1340                1345                1350

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr
    1355                1360                1365

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Thr  Ser Thr Glu
    1370                1375                1380

Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1385                1390                1395

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1400                1405                1410

Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Ser
    1415                1420                1425

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    1430                1435                1440

Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu
    1445                1450                1455

Gly Ser  Ala Pro Gly Ala Ser  Ser Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485
```

```
Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
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Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                 1510                 1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                 1525                 1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                 1540                 1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                 1555                 1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                 1570                 1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                 1585                 1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                 1600                 1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                 1615                 1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                 1630                 1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                 1645                 1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                 1660                 1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                 1675                 1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                 1690                 1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                 1705                 1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                 1720                 1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                 1735                 1740
```

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Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                1960                1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                1975                1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                1990                1995
```

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2000 2005 2010

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
2015 2020 2025

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
2030 2035 2040

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2045 2050 2055

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
2060 2065 2070

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
2075 2080 2085

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
2090 2095 2100

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
2105 2110 2115

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
2120 2125 2130

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
2135 2140 2145

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2150 2155 2160

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2165 2170 2175

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
2180 2185 2190

Pro Ala Thr Gly His His His His His His
2195 2200

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<211> 6612
<212> DNA
<213> Artificial Sequence

<220>
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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

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catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

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atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

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acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

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cctgggacat ccaccgagcc ctccgaagga tcagccccag gcaccagcac ggagccctct 1560

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<211> 2203
<212> PRT
<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1329

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu

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                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
                405                 410                 415

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser
            420                 425                 430

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly
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435 440 445

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
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Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala
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Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly
485 490 495

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
500 505 510

Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala
515 520 525

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
530 535 540

Ser Ser Thr Gly Ser Pro Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545 550 555 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
565 570 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
580 585 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
595 600 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
610 615 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
645 650 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys

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705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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980 985 990

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
995 1000 1005

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser
1010 1015 1020

Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
1025 1030 1035

Ser Ala Pro Gly Ala Ser Ser Pro Pro Val Leu Lys Arg His Gln
1040 1045 1050

Ala Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
1055 1060 1065

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
1070 1075 1080

Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln
1085 1090 1095

Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp
1100 1105 1110

Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala
1115 1120 1125

Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu
1130 1135 1140

Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
1145 1150 1155

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1160 1165 1170

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1175 1180 1185

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1190 1195 1200

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1205 1210 1215

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1220 1225 1230

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp

1235 1240 1245

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1250 1255 1260

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1265 1270 1275

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1280 1285 1290

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
1295 1300 1305

Cys Asn Ile Gln Met Glu Asp Pro Gly Ala Pro Ser Pro Ala Gly
1310 1315 1320

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
1325 1330 1335

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
1340 1345 1350

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr
1355 1360 1365

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
1370 1375 1380

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
1385 1390 1395

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1400 1405 1410

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser
1415 1420 1425

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
1430 1435 1440

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
1445 1450 1455

Gly Ser Ala Pro Gly Ala Ser Ser Thr Phe Lys Glu Asn Tyr Arg
1460 1465 1470

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1475 1480 1485

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met

1490 1495 1500

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1505 1510 1515

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1520 1525 1530

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1535 1540 1545

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1550 1555 1560

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1565 1570 1575

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1580 1585 1590

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1595 1600 1605

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1610 1615 1620

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1625 1630 1635

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1640 1645 1650

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1655 1660 1665

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1670 1675 1680

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1685 1690 1695

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1700 1705 1710

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1715 1720 1725

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1730 1735 1740

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr

1745 1750 1755

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1760 1765 1770

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1775 1780 1785

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1790 1795 1800

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1805 1810 1815

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1820 1825 1830

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1835 1840 1845

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1850 1855 1860

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1865 1870 1875

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Pro Gly Thr
1880 1885 1890

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1895 1900 1905

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
1910 1915 1920

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1925 1930 1935

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1940 1945 1950

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
1955 1960 1965

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
1970 1975 1980

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1985 1990 1995

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser

2000 2005 2010

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
2015 2020 2025

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
2030 2035 2040

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2045 2050 2055

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
2060 2065 2070

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
2075 2080 2085

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
2090 2095 2100

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
2105 2110 2115

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
2120 2125 2130

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
2135 2140 2145

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2150 2155 2160

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2165 2170 2175

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
2180 2185 2190

Pro Ala Thr Gly His His His His His His
2195 2200

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<211> 6612
<212> DNA
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<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

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tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

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atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

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tcgagcgaac aaaaactcat ctcagaagag gatctgtcac ctgcaaccgg tcatcatcac 6600

catcaccatt ga 6612

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<211> 2203
<212> PRT
<213> Artificial Sequence

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

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Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
                405                 410                 415

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            420                 425                 430

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
        435                 440                 445
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Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    450                 455                 460

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
465                 470                 475                 480

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
                485                 490                 495

Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            500                 505                 510

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
        515                 520                 525

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
    530                 535                 540

Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720
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Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990
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Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
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Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245
```

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Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Gly Ala Pro Gly  Thr Pro Gly
    1310                1315                1320

Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1325                1330                1335

Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1340                1345                1350

Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly
    1355                1360                1365

Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly Pro Gly  Thr Pro Gly
    1370                1375                1380

Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1385                1390                1395

Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1400                1405                1410

Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly
    1415                1420                1425

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr
    1430                1435                1440

Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1445                1450                1455

Ser Thr  Gly Ser Pro Ala Ser  Ser Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500
```

```
Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                1555                1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                1615                1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755
```

```
Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                1960                1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                1975                1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                1990                1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                2005                2010
```

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
2015 2020 2025

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
2030 2035 2040

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2045 2050 2055

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
2060 2065 2070

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
2075 2080 2085

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
2090 2095 2100

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
2105 2110 2115

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
2120 2125 2130

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
2135 2140 2145

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2150 2155 2160

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2165 2170 2175

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Ala Thr Gly His His His His His His
2195 2200

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<211> 6612
<212> DNA
<213> Artificial Sequence

<220>
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gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa      360
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ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct      480
catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga      540
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tataaatgga cagtgactgt agaagatggg ccaactaaat cagatcctcg gtgcctgacc     2040
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<211> 2203
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1333

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

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Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
                405                 410                 415

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser
            420                 425                 430

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly
        435                 440                 445
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Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
    450                 455                 460

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala
465                 470                 475                 480

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly
                485                 490                 495

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
            500                 505                 510

Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala
        515                 520                 525

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
    530                 535                 540

Ser Ser Thr Gly Ser Pro Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720
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Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990
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Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
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Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
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Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245
```

```
Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
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Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Gly Ala Pro Gly  Thr Pro Gly
    1310                1315                1320

Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1325                1330                1335

Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1340                1345                1350

Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly
    1355                1360                1365

Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly Pro Gly  Thr Pro Gly
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Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
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Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
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Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly
    1415                1420                1425

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr
    1430                1435                1440

Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1445                1450                1455

Ser Thr  Gly Ser Pro Ala Ser  Ser Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500
```

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Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
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Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                1555                1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
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Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755
```

```
Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                1960                1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                1975                1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                1990                1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                2005                2010
```

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
2015 2020 2025

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
2030 2035 2040

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2045 2050 2055

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
2060 2065 2070

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
2075 2080 2085

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
2090 2095 2100

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
2105 2110 2115

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
2120 2125 2130

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
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Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
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Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Ala Thr Gly His His His His His His
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atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

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Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445
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    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720
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Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
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Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
        755                 760                 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
    770                 775                 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785                 790                 795                 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                805                 810                 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            820                 825                 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
        835                 840                 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
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Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865                 870                 875                 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
            980                 985                 990
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Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
995 1000 1005

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
1010 1015 1020

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
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Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
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Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
1055 1060 1065

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
1070 1075 1080

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
1085 1090 1095

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
1100 1105 1110

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser
1115 1120 1125

Ser Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val
1130 1135 1140

Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg
1145 1150 1155

Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
1160 1165 1170

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala
1175 1180 1185

Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu
1190 1195 1200

Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro
1205 1210 1215

Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala
1220 1225 1230

Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser
1235 1240 1245

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Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
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Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
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Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1280                1285                1290

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1295                1300                1305

Arg Gly  Ala Pro Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1310                1315                1320

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1325                1330                1335

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1340                1345                1350

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1355                1360                1365

Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1370                1375                1380

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    1385                1390                1395

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Ser Glu  Pro Ala Thr
    1400                1405                1410

Ser Gly  Ser Glu Thr Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1415                1420                1425

Thr Glu  Glu Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1430                1435                1440

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Ala Ser
    1445                1450                1455

Ser Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1460                1465                1470

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1475                1480                1485

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1490                1495                1500
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Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
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Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1520                1525                1530

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1535                1540                1545

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1550                1555                1560

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1565                1570                1575

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1580                1585                1590

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1595                1600                1605

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1610                1615                1620

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1625                1630                1635

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1640                1645                1650

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1655                1660                1665

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1670                1675                1680

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1685                1690                1695

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1700                1705                1710

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1715                1720                1725

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1730                1735                1740

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1745                1750                1755
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Phe Ala Thr Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly
1760 1765 1770

Arg Ser Asn Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp
1775 1780 1785

Leu Gln Val Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr
1790 1795 1800

Thr Gln Gly Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu
1805 1810 1815

Phe Leu Ile Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe
1820 1825 1830

Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser
1835 1840 1845

Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg
1850 1855 1860

Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu
1865 1870 1875

Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala
1880 1885 1890

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
1895 1900 1905

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
1910 1915 1920

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
1925 1930 1935

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1940 1945 1950

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
1955 1960 1965

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
1970 1975 1980

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
1985 1990 1995

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2000 2005 2010

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser
2015 2020 2025

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu
2030 2035 2040

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
2045 2050 2055

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2060 2065 2070

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2075 2080 2085

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala
2090 2095 2100

Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
2105 2110 2115

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2120 2125 2130

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
2135 2140 2145

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
2150 2155 2160

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
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Gly Ser Ala Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu
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Asp Leu Ser Pro Ala Thr Gly His His His His His His
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<211> 6621
<212> DNA
<213> Artificial Sequence

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atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

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catcatcacc atcaccattg a 6621

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<212> PRT
<213> Artificial Sequence

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20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu

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            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
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    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
```

725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
740 745 750

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
755 760 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
770 775 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785 790 795 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
805 810 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
820 825 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
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Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
850 855 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865 870 875 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
885 890 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
900 905 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
915 920 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
930 935 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945 950 955 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
965 970 975

Arg Ala Gln Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
980 985 990

Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala

995 1000 1005

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
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Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
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Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
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Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
1055 1060 1065

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
1070 1075 1080

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
1085 1090 1095

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
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Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser
1115 1120 1125

Ser Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val
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Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg
1145 1150 1155

Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
1160 1165 1170

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala
1175 1180 1185

Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu
1190 1195 1200

Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro
1205 1210 1215

Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala
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Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser
1235 1240 1245

Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro

1250 1255 1260

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg
1265 1270 1275

Gln Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp
1280 1285 1290

Glu Thr Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys
1295 1300 1305

Arg Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
1310 1315 1320

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
1325 1330 1335

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
1340 1345 1350

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
1355 1360 1365

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
1370 1375 1380

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
1385 1390 1395

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
1400 1405 1410

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
1415 1420 1425

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
1430 1435 1440

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser
1445 1450 1455

Ser Ala Pro Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu
1460 1465 1470

Asn Tyr Arg Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu
1475 1480 1485

Pro Gly Leu Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu
1490 1495 1500

Leu Ser Met Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser

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Gly His Val Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala
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Leu Tyr Asn Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu
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Pro Ser Lys Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu
1550 1555 1560

His Leu His Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn
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Lys Cys Gln Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp
1580 1585 1590

Phe Gln Ile Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys
1595 1600 1605

Leu Ala Arg Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr
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Lys Glu Pro Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met
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Ile Ile His Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser
1640 1645 1650

Ser Leu Tyr Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly
1655 1660 1665

Lys Lys Trp Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met
1670 1675 1680

Val Phe Phe Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile
1685 1690 1695

Phe Asn Pro Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr
1700 1705 1710

His Tyr Ser Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys
1715 1720 1725

Asp Leu Asn Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala
1730 1735 1740

Ile Ser Asp Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met
1745 1750 1755

Phe Ala Thr Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly

1760 1765 1770

Arg Ser Asn Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp
1775 1780 1785

Leu Gln Val Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr
1790 1795 1800

Thr Gln Gly Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu
1805 1810 1815

Phe Leu Ile Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe
1820 1825 1830

Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser
1835 1840 1845

Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg
1850 1855 1860

Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu
1865 1870 1875

Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala
1880 1885 1890

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
1895 1900 1905

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
1910 1915 1920

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
1925 1930 1935

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1940 1945 1950

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
1955 1960 1965

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
1970 1975 1980

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
1985 1990 1995

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2000 2005 2010

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser

2015 2020 2025

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu
2030 2035 2040

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
2045 2050 2055

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2060 2065 2070

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2075 2080 2085

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala
2090 2095 2100

Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
2105 2110 2115

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2120 2125 2130

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
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Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
2150 2155 2160

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
2165 2170 2175

Gly Ser Ala Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu
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Asp Leu Ser Pro Ala Thr Gly His His His His His His
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<211> 6621
<212> DNA
<213> Artificial Sequence

<220>
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<212> PRT
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

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His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460
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Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735
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Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
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Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
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Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
    770                 775                 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785                 790                 795                 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                805                 810                 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            820                 825                 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
        835                 840                 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    850                 855                 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865                 870                 875                 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
            980                 985                 990

Ser Pro Gly Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
        995                 1000                1005
```

```
Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
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Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
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Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1040                 1045                 1050

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1055                 1060                 1065

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1070                 1075                 1080

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1085                 1090                 1095

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1100                 1105                 1110

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
    1115                 1120                 1125

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1130                 1135                 1140

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1145                 1150                 1155

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1160                 1165                 1170

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1175                 1180                 1185

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1190                 1195                 1200

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1205                 1210                 1215

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1220                 1225                 1230

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1235                 1240                 1245

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1250                 1255                 1260
```

```
Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
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Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1280                1285                1290

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1295                1300                1305

Arg Ala  Pro Cys Asn Ile Gly  Ala Pro Ser Pro Ala  Gly Ser Pro
    1310                1315                1320

Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1325                1330                1335

Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    1340                1345                1350

Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr
    1355                1360                1365

Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser
    1370                1375                1380

Glu Gly  Ser Ala Pro Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1385                1390                1395

Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly
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Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr Pro Gly  Ser Pro Ala
    1415                1420                1425

Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr Ser Glu  Ser Ala Thr
    1430                1435                1440

Pro Glu  Ser Gly Pro Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    1445                1450                1455

Ala Pro  Gly Ala Ser Ser Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1460                1465                1470

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1475                1480                1485

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1490                1495                1500

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1505                1510                1515
```

```
Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1520                1525                1530

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1535                1540                1545

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1550                1555                1560

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1565                1570                1575

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1580                1585                1590

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1595                1600                1605

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1610                1615                1620

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1625                1630                1635

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1640                1645                1650

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1655                1660                1665

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1670                1675                1680

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1685                1690                1695

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1700                1705                1710

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1715                1720                1725

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1730                1735                1740

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1745                1750                1755

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1760                1765                1770
```

```
Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
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Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1790                 1795                 1800

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1805                 1810                 1815

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1820                 1825                 1830

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1835                 1840                 1845

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    1850                 1855                 1860

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    1865                 1870                 1875

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    1880                 1885                 1890

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1895                 1900                 1905

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    1910                 1915                 1920

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1925                 1930                 1935

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1940                 1945                 1950

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    1955                 1960                 1965

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    1970                 1975                 1980

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1985                 1990                 1995

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2000                 2005                 2010

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser
    2015                 2020                 2025
```

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu
2030 2035 2040

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
2045 2050 2055

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2060 2065 2070

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2075 2080 2085

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala
2090 2095 2100

Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
2105 2110 2115

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
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Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
2150 2155 2160

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
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Gly Ser Ala Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu
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Asp Leu Ser Pro Ala Thr Gly His His His His His His
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

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His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460
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Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735
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Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
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Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
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Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
    770                 775                 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785                 790                 795                 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                805                 810                 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            820                 825                 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
        835                 840                 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    850                 855                 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865                 870                 875                 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
            980                 985                 990

Ser Pro Gly Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
        995                 1000                1005
```

```
Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
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Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
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Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1040                 1045                 1050

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1055                 1060                 1065

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1070                 1075                 1080

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1085                 1090                 1095

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1100                 1105                 1110

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
    1115                 1120                 1125

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1130                 1135                 1140

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1145                 1150                 1155

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1160                 1165                 1170

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1175                 1180                 1185

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1190                 1195                 1200

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1205                 1210                 1215

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1220                 1225                 1230

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1235                 1240                 1245

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1250                 1255                 1260
```

```
Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
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Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1280                1285                1290

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1295                1300                1305

Arg Ala  Pro Cys Asn Ile Gly  Ala Pro Gly Thr Pro  Gly Ser Gly
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Thr Ala  Ser Ser Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
    1325                1330                1335

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
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Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser
    1355                1360                1365

Pro Ser  Ala Ser Thr Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly
    1370                1375                1380

Thr Ala  Ser Ser Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
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Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1400                1405                1410

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser
    1415                1420                1425

Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser
    1430                1435                1440

Gly Ala  Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
    1445                1450                1455

Gly Ser  Pro Ala Ser Ser Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1460                1465                1470

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1475                1480                1485

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1490                1495                1500

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1505                1510                1515
```

```
Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
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Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1535                1540                1545

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1550                1555                1560

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1565                1570                1575

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1580                1585                1590

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1595                1600                1605

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1610                1615                1620

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1625                1630                1635

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1640                1645                1650

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1655                1660                1665

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1670                1675                1680

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1685                1690                1695

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1700                1705                1710

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1715                1720                1725

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1730                1735                1740

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1745                1750                1755

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
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```

```
Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
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Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1790                 1795                 1800

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1805                 1810                 1815

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1820                 1825                 1830

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1835                 1840                 1845

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    1850                 1855                 1860

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    1865                 1870                 1875

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    1880                 1885                 1890

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1895                 1900                 1905

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    1910                 1915                 1920

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1925                 1930                 1935

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1940                 1945                 1950

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    1955                 1960                 1965

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    1970                 1975                 1980

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1985                 1990                 1995

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2000                 2005                 2010

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser
    2015                 2020                 2025
```

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu
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Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
2045 2050 2055

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2060 2065 2070

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2075 2080 2085

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala
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Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
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Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
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Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
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Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
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Gly Ser Ala Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu
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Asp Leu Ser Pro Ala Thr Gly His His His His His His
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

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His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
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Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
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Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460
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Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735
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Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
            740                 745                 750

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
        755                 760                 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
    770                 775                 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785                 790                 795                 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                805                 810                 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            820                 825                 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
        835                 840                 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    850                 855                 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865                 870                 875                 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
            980                 985                 990

Ser Pro Gly Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
        995                 1000                1005
```

```
Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
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Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
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Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1040                 1045                 1050

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1055                 1060                 1065

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1070                 1075                 1080

Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1085                 1090                 1095

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1100                 1105                 1110

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Ala Ser
    1115                 1120                 1125

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1130                 1135                 1140

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1145                 1150                 1155

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1160                 1165                 1170

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1175                 1180                 1185

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1190                 1195                 1200

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1205                 1210                 1215

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1220                 1225                 1230

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1235                 1240                 1245

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1250                 1255                 1260
```

```
Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
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Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1280                1285                1290

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1295                1300                1305

Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Gly  Ala Pro Ser
    1310                1315                1320

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    1325                1330                1335

Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu
    1340                1345                1350

Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    1355                1360                1365

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
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Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Thr  Ser Glu Ser
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Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
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Ser Glu  Thr Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
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Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr
    1430                1435                1440

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Thr  Ser Thr Glu
    1445                1450                1455

Pro Ser  Glu Gly Ser Ala Pro  Gly Ala Ser Ser Thr  Phe Lys Glu
    1460                1465                1470

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1475                1480                1485

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1490                1495                1500

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1505                1510                1515
```

```
Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1520                1525                1530

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1535                1540                1545

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1550                1555                1560

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1565                1570                1575

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1580                1585                1590

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1595                1600                1605

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1610                1615                1620

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1625                1630                1635

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1640                1645                1650

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1655                1660                1665

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1670                1675                1680

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1685                1690                1695

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1700                1705                1710

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1715                1720                1725

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1730                1735                1740

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1745                1750                1755

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1760                1765                1770
```

```
Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1775                 1780                 1785

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1790                 1795                 1800

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1805                 1810                 1815

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1820                 1825                 1830

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1835                 1840                 1845

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    1850                 1855                 1860

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    1865                 1870                 1875

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    1880                 1885                 1890

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1895                 1900                 1905

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    1910                 1915                 1920

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1925                 1930                 1935

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1940                 1945                 1950

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    1955                 1960                 1965

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    1970                 1975                 1980

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1985                 1990                 1995

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2000                 2005                 2010

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser
    2015                 2020                 2025
```

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu
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Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
2045 2050 2055

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2060 2065 2070

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2075 2080 2085

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala
2090 2095 2100

Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
2105 2110 2115

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2120 2125 2130

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
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Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
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Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
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Asp Leu Ser Pro Ala Thr Gly His His His His His His
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp

195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
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Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
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Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
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Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile

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465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
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His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
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Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
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Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
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Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
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Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
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Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
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Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
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Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
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Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
755 760 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
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Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785 790 795 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
805 810 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
820 825 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
835 840 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
850 855 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865 870 875 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
885 890 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
900 905 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
915 920 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
930 935 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945 950 955 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
965 970 975

Arg Ala Gln Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
980 985 990

Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
995 1000 1005

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly

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Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
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Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
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Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
1055 1060 1065

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
1070 1075 1080

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
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Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
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Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser
1115 1120 1125

Ser Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val
1130 1135 1140

Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg
1145 1150 1155

Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
1160 1165 1170

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala
1175 1180 1185

Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu
1190 1195 1200

Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro
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Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala
1220 1225 1230

Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser
1235 1240 1245

Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
1250 1255 1260

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg

1265 1270 1275

Gln Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp
1280 1285 1290

Glu Thr Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys
1295 1300 1305

Arg Ala Pro Cys Asn Ile Gln Met Glu Asp Pro Gly Ala Pro Gly
1310 1315 1320

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
1325 1330 1335

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
1340 1345 1350

Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
1355 1360 1365

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
1370 1375 1380

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
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Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
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Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
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Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
1430 1435 1440

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro
1445 1450 1455

Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Thr Phe Lys Glu
1460 1465 1470

Asn Tyr Arg Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu
1475 1480 1485

Pro Gly Leu Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu
1490 1495 1500

Leu Ser Met Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser
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Gly His Val Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala

1520 1525 1530

Leu Tyr Asn Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu
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Pro Ser Lys Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu
1550 1555 1560

His Leu His Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn
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Lys Cys Gln Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp
1580 1585 1590

Phe Gln Ile Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys
1595 1600 1605

Leu Ala Arg Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr
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Lys Glu Pro Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met
1625 1630 1635

Ile Ile His Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser
1640 1645 1650

Ser Leu Tyr Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly
1655 1660 1665

Lys Lys Trp Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met
1670 1675 1680

Val Phe Phe Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile
1685 1690 1695

Phe Asn Pro Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr
1700 1705 1710

His Tyr Ser Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys
1715 1720 1725

Asp Leu Asn Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala
1730 1735 1740

Ile Ser Asp Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met
1745 1750 1755

Phe Ala Thr Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly
1760 1765 1770

Arg Ser Asn Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp

1775 1780 1785

Leu Gln Val Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr
1790 1795 1800

Thr Gln Gly Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu
1805 1810 1815

Phe Leu Ile Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe
1820 1825 1830

Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser
1835 1840 1845

Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg
1850 1855 1860

Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu
1865 1870 1875

Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala
1880 1885 1890

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
1895 1900 1905

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
1910 1915 1920

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
1925 1930 1935

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1940 1945 1950

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
1955 1960 1965

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
1970 1975 1980

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
1985 1990 1995

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2000 2005 2010

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser
2015 2020 2025

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu

2030 2035 2040

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
2045 2050 2055

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2060 2065 2070

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2075 2080 2085

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala
2090 2095 2100

Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
2105 2110 2115

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2120 2125 2130

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
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Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
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Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
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Asp Leu Ser Pro Ala Thr Gly His His His His His His
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<211> 6621
<212> DNA
<213> Artificial Sequence

<220>
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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
35 40 45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
50 55 60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65 70 75 80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
85 90 95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
100 105 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
115 120 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
130 135 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145 150 155 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

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Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480
```

```
Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Gly Ala Pro Gly Ala Ser Pro
545                 550                 555                 560

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
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Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            580                 585                 590

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
        595                 600                 605

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
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Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
625                 630                 635                 640

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
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Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
            660                 665                 670

Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
        675                 680                 685

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Asp
    690                 695                 700

Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750
```

```
Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
        755                 760                 765

Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
    1010                1015                1020
```

```
Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
    1025                1030                1035

Pro Arg  Ser Phe Ser Gln Asn  Gly Ala Pro Ser Pro  Ala Gly Ser
    1040                1045                1050

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1055                1060                1065

Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro
    1070                1075                1080

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1085                1090                1095

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Thr Glu Pro
    1100                1105                1110

Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1115                1120                1125

Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1130                1135                1140

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1145                1150                1155

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1160                1165                1170

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1175                1180                1185

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1190                1195                1200

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1205                1210                1215

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1220                1225                1230

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1235                1240                1245

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1250                1255                1260

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1265                1270                1275
```

```
Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1280                1285                1290

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1295                1300                1305

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1310                1315                1320

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1325                1330                1335

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1340                1345                1350

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1355                1360                1365

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1370                1375                1380

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1385                1390                1395

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1400                1405                1410

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1415                1420                1425

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1430                1435                1440

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1445                1450                1455

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530
```

```
Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                1555                1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                1615                1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
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Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
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Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785
```

```
Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
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Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                1960                1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                1975                1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                1990                1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                2005                2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                2020                2025

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    2030                2035                2040
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Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
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Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
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Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
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Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
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Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
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Pro Ala Thr Gly His His His His His His
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        35                  40                  45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
    50                  55                  60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65                  70                  75                  80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
                85                  90                  95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
            100                 105                 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
        115                 120                 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
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Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145                 150                 155                 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205
```

```
Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480
```

```
Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
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Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
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Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gly
545                 550                 555                 560

Ala Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser
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Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala
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Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
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Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser
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Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr
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Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
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Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr
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Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr
        675                 680                 685

Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
    690                 695                 700

Ser Pro Ala Ser Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly
705                 710                 715                 720

Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe
                725                 730                 735

Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu
            740                 745                 750
```

```
Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln
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Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg
    770                 775                 780

Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp
785                 790                 795                 800

Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr
                805                 810                 815

Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr
            820                 825                 830

Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
        835                 840                 845

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn
    850                 855                 860

Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp
865                 870                 875                 880

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
                885                 890                 895

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
            900                 905                 910

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser
        915                 920                 925

Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala
    930                 935                 940

Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His
945                 950                 955                 960

Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu
                965                 970                 975

Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
            980                 985                 990

His Asn Ser Asp Phe Arg Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val
        995                 1000                1005

Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr
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```

```
Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu
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Pro Arg  Ser Phe Ser Gln Asn  Gly Ala Pro Ser Pro  Ala Gly Ser
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Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1055                1060                1065

Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro
    1070                1075                1080

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1085                1090                1095

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Thr Glu Pro
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Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
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Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
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Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1145                1150                1155

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1160                1165                1170

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1175                1180                1185

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
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Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1205                1210                1215

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1220                1225                1230

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1235                1240                1245

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1250                1255                1260

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1265                1270                1275
```

```
Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
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Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1295                1300                1305

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1310                1315                1320

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
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Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1340                1345                1350

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
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Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1370                1375                1380

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1385                1390                1395

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1400                1405                1410

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1415                1420                1425

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1430                1435                1440

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1445                1450                1455

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530
```

```
Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
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Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
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Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
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Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
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Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
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Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
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Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
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Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
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Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
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Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
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Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
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Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
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Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
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Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
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Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
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Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
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Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
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Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
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Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                1960                1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                1975                1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                1990                1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                2005                2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
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Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    2030                2035                2040
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Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
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Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
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Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
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tcagcacccg gctctgagcc ggccacaagt ggcagtgaga cacccggcac ttcagagagt 6480

gccacccccg agagtggccc aggcactagt accgagccct ctgaaggcag tgcgccagcc 6540

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catcaccatt ga 6612

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<211> 2203
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Gly Ala Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
35 40 45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
50 55 60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65 70 75 80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
85 90 95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
100 105 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
115 120 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
130 135 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145 150 155 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
165 170 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
180 185 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
195 200 205

```
Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480
```

```
Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750
```

```
Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020
```

```
Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275
```

```
Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1295                1300                1305

Pro Ser  Pro Ala Gly Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser
    1310                1315                1320

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro
    1325                1330                1335

Ser Glu  Gly Ser Ala Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1340                1345                1350

Thr Glu  Glu Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1355                1360                1365

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1370                1375                1380

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
    1385                1390                1395

Ser Gly  Ser Glu Thr Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1400                1405                1410

Glu Thr  Pro Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1415                1420                1425

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1430                1435                1440

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ala Ser  Ser Ala Pro
    1445                1450                1455

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530
```

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1535 1540 1545

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1550 1555 1560

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1565 1570 1575

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1580 1585 1590

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1595 1600 1605

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1610 1615 1620

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1625 1630 1635

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1640 1645 1650

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1655 1660 1665

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1670 1675 1680

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1685 1690 1695

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1700 1705 1710

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1715 1720 1725

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1730 1735 1740

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1745 1750 1755

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1760 1765 1770

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1775 1780 1785

```
Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                1855                1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                1870                1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1895                1900                1905

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1910                1915                1920

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                1960                1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                1975                1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                1990                1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                2005                2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                2020                2025

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    2030                2035                2040
```

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2045 2050 2055

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
2060 2065 2070

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
2075 2080 2085

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
2090 2095 2100

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
2105 2110 2115

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
2120 2125 2130

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
2135 2140 2145

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2150 2155 2160

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2165 2170 2175

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
2180 2185 2190

Pro Ala Thr Gly His His His His His His
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<211> 2203
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1353

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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
        35                  40                  45

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
    50                  55                  60

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
65                  70                  75                  80

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
                85                  90                  95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
            100                 105                 110

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
        115                 120                 125

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
    130                 135                 140

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145                 150                 155                 160

Ser Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
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210 215 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225 230 235 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
245 250 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
260 265 270

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
275 280 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
290 295 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305 310 315 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
325 330 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
340 345 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
355 360 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
370 375 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385 390 395 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
405 410 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
420 425 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
435 440 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
450 455 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465 470 475 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr

485 490 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
500 505 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
515 520 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
530 535 540

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545 550 555 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
565 570 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
580 585 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
595 600 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
610 615 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625 630 635 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
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Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
660 665 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
675 680 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
690 695 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705 710 715 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
725 730 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
740 745 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn

```
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Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
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Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
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Ser Ala Pro Gly Ala Ser Ser Pro Pro Val Leu Lys Arg His Gln
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Ala Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
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Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
1070 1075 1080

Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln
1085 1090 1095

Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp
1100 1105 1110

Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala
1115 1120 1125

Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu
1130 1135 1140

Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
1145 1150 1155

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1160 1165 1170

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1175 1180 1185

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1190 1195 1200

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1205 1210 1215

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1220 1225 1230

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1235 1240 1245

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1250 1255 1260

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1265 1270 1275

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys

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Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
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Cys Asn  Ile Gly Ala Pro Ser  Pro Ala Gly Ser Pro  Thr Ser Thr
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Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
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Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala Pro Gly  Ser Pro Ala
    1340                1345                1350

Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr Ser Thr  Glu Pro Ser
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Glu Gly  Ser Ala Pro Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    1370                1375                1380

Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    1385                1390                1395

Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr Pro Gly  Ser Glu Pro
    1400                1405                1410

Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Ser Pro Ala  Gly Ser Pro
    1415                1420                1425

Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1430                1435                1440

Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    1445                1450                1455

Ala Ser  Ser Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
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Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
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Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1565 1570 1575

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1580 1585 1590

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1595 1600 1605

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1610 1615 1620

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
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Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1640 1645 1650

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1655 1660 1665

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1670 1675 1680

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1685 1690 1695

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1700 1705 1710

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1715 1720 1725

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1730 1735 1740

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1745 1750 1755

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1760 1765 1770

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1775 1780 1785

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly

1790 1795 1800

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1805 1810 1815

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1820 1825 1830

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1835 1840 1845

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1850 1855 1860

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1865 1870 1875

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Pro Gly Thr
1880 1885 1890

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1895 1900 1905

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
1910 1915 1920

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1925 1930 1935

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1940 1945 1950

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
1955 1960 1965

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
1970 1975 1980

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1985 1990 1995

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2000 2005 2010

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
2015 2020 2025

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
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Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly

2045 2050 2055

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
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Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
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Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
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Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
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Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
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Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
2135 2140 2145

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
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Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

```
Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
                405                 410                 415

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            420                 425                 430

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
        435                 440                 445

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    450                 455                 460

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
465                 470                 475                 480

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
                485                 490                 495
```

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Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            500                 505                 510

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
        515                 520                 525

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
    530                 535                 540

Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765
```

```
Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035
```

```
Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Gly  Ala Pro Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1130                1135                1140

Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser
    1145                1150                1155

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
    1160                1165                1170

Ser Gly  Ser Glu Thr Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1175                1180                1185

Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1190                1195                1200

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Pro
    1205                1210                1215

Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Ser Pro  Ala Gly Ser
    1220                1225                1230

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1235                1240                1245

Thr Glu  Glu Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1250                1255                1260

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Ala Ser
    1265                1270                1275

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1280                1285                1290
```

```
Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
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Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1310                1315                1320

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1325                1330                1335

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1340                1345                1350

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1355                1360                1365

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1370                1375                1380

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1385                1390                1395

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1400                1405                1410

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1415                1420                1425

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1430                1435                1440

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1445                1450                1455

Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1460                1465                1470

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1475                1480                1485

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1490                1495                1500

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1505                1510                1515

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1520                1525                1530

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1535                1540                1545
```

```
Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1550                1555                1560

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1565                1570                1575

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1580                1585                1590

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1595                1600                1605

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1610                1615                1620

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1625                1630                1635

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1640                1645                1650

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1655                1660                1665

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1670                1675                1680

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1685                1690                1695

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1700                1705                1710

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1715                1720                1725

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1730                1735                1740

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1745                1750                1755

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1760                1765                1770

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1775                1780                1785

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1790                1795                1800
```

```
Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1805                1810                1815

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1820                1825                1830

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1835                1840                1845

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    1850                1855                1860

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    1865                1870                1875

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    1880                1885                1890

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1895                1900                1905

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    1910                1915                1920

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1925                1930                1935

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1940                1945                1950

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    1955                1960                1965

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    1970                1975                1980

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1985                1990                1995

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2000                2005                2010

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser
    2015                2020                2025

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Thr Glu
    2030                2035                2040

Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2045                2050                2055
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Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2060 2065 2070

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2075 2080 2085

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala
2090 2095 2100

Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
2105 2110 2115

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2120 2125 2130

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
2135 2140 2145

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
2150 2155 2160

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
2165 2170 2175

Gly Ser Ala Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu
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Asp Leu Ser Pro Ala Thr Gly His His His His His His
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220
```

```
Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gly Ala Pro Thr Ser Thr Glu
                405                 410                 415

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
            420                 425                 430

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
        435                 440                 445

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
    450                 455                 460

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
465                 470                 475                 480

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
                485                 490                 495
```

```
Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu
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Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
        515                 520                 525

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
    530                 535                 540

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ala Ser Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765
```

```
Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035
```

```
Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095

Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                1105                1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290
```

```
Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
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Cys Asn  Ile Gln Met Glu Asp  Pro Gly Ala Pro Gly  Thr Pro Gly
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Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1325                1330                1335

Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1340                1345                1350

Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly
    1355                1360                1365

Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly Pro Gly  Thr Pro Gly
    1370                1375                1380

Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1385                1390                1395

Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1400                1405                1410

Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly
    1415                1420                1425

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr
    1430                1435                1440

Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1445                1450                1455

Ser Thr  Gly Ser Pro Ala Ser  Ser Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545
```

```
Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
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Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                1615                1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800
```

```
Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
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Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                 1825                 1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                 1840                 1845

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1850                 1855                 1860

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1865                 1870                 1875

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1880                 1885                 1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
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Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
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Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1925                 1930                 1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                 1945                 1950

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1955                 1960                 1965

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1970                 1975                 1980

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                 1990                 1995

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2000                 2005                 2010

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2015                 2020                 2025

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    2030                 2035                 2040

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2045                 2050                 2055
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Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
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Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
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Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
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Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
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Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
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Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
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Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2150 2155 2160

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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<223> Description of Artificial Sequence: Synthetic polypeptide

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Gly Ala Pro Gly
            180                 185                 190

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
        195                 200                 205

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
    210                 215                 220
```

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Glu Thr Pro Ala Ser Ser Thr Gln Thr Leu His Lys Phe Ile Leu Leu
225                 230                 235                 240

Phe Ala Val Phe Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn
                245                 250                 255

Ser Leu Met Gln Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys
            260                 265                 270

Met His Thr Val Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile
        275                 280                 285

Gly Cys His Arg Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr
    290                 295                 300

Thr Pro Glu Val His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val
305                 310                 315                 320

Arg Asn His Arg Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu
                325                 330                 335

Thr Ala Gln Thr Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys
            340                 345                 350

His Ile Ser Ser His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val
        355                 360                 365

Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu
    370                 375                 380

Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val
385                 390                 395                 400

Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val
                405                 410                 415

Ala Lys Lys His Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu
            420                 425                 430

Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser
        435                 440                 445

Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys
    450                 455                 460

Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr
465                 470                 475                 480

Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr
                485                 490                 495
```

```
Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser
            500                 505                 510

Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu
        515                 520                 525

Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro
    530                 535                 540

Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu
545                 550                 555                 560

Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser
                565                 570                 575

Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro
            580                 585                 590

Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile
        595                 600                 605

Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn
    610                 615                 620

Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro
625                 630                 635                 640

Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met
                645                 650                 655

His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys
            660                 665                 670

Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr
        675                 680                 685

Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met
    690                 695                 700

Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val
705                 710                 715                 720

Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn
                725                 730                 735

Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser
            740                 745                 750

Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile
        755                 760                 765
```

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Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe
    770                 775                 780

Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
785                 790                 795                 800

Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
                805                 810                 815

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
            820                 825                 830

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
        835                 840                 845

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
    850                 855                 860

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
865                 870                 875                 880

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
                885                 890                 895

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro
            900                 905                 910

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
        915                 920                 925

Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro Pro Val Leu Lys Arg His
    930                 935                 940

Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
945                 950                 955                 960

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
                965                 970                 975

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
            980                 985                 990

Thr Arg His Tyr Phe Ile Ala Ala  Val Glu Arg Leu Trp  Asp Tyr Gly
        995                 1000                1005

Met Ser  Ser Ser Pro His Val  Leu Arg Asn Arg Ala  Gln Ser Gly
    1010                1015                1020

Ser Val  Pro Gln Phe Lys Lys  Val Val Phe Gln Glu  Phe Thr Asp
    1025                1030                1035
```

```
Gly Ser  Phe Thr Gln Pro Leu  Tyr Arg Gly Glu Leu  Asn Glu His
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Leu Gly  Leu Leu Gly Pro Tyr  Ile Arg Ala Glu Val  Glu Asp Asn
    1055                1060                1065

Ile Met  Val Thr Phe Arg Asn  Gln Ala Ser Arg Pro  Tyr Ser Phe
    1070                1075                1080

Tyr Ser  Ser Leu Ile Ser Tyr  Glu Glu Asp Gln Arg  Gln Gly Ala
    1085                1090                1095

Glu Pro  Arg Lys Asn Phe Val  Lys Pro Asn Glu Thr  Lys Thr Tyr
    1100                1105                1110

Phe Trp  Lys Val Gln His His  Met Ala Pro Thr Lys  Asp Glu Phe
    1115                1120                1125

Asp Cys  Lys Ala Trp Ala Tyr  Phe Ser Asp Val Asp  Leu Glu Lys
    1130                1135                1140

Asp Val  His Ser Gly Leu Ile  Gly Pro Leu Leu Val  Cys His Thr
    1145                1150                1155

Asn Thr  Leu Asn Pro Ala His  Gly Arg Gln Val Thr  Val Gln Glu
    1160                1165                1170

Phe Ala  Leu Phe Phe Thr Ile  Phe Asp Glu Thr Lys  Ser Trp Tyr
    1175                1180                1185

Phe Thr  Glu Asn Met Glu Arg  Asn Cys Arg Gly Ala  Pro Thr Ser
    1190                1195                1200

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
    1205                1210                1215

Ser Gly  Ser Glu Thr Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1220                1225                1230

Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro
    1235                1240                1245

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1250                1255                1260

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala
    1265                1270                1275

Thr Pro  Glu Ser Gly Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1280                1285                1290
```

```
Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
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Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser
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Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro
    1325                 1330                 1335

Ser Glu  Gly Ser Ala Pro Gly  Ala Ser Ser Ala Pro  Cys Asn Ile
    1340                 1345                 1350

Gln Met  Glu Asp Pro Thr Phe  Lys Glu Asn Tyr Arg  Phe His Ala
    1355                 1360                 1365

Ile Asn  Gly Tyr Ile Met Asp  Thr Leu Pro Gly Leu  Val Met Ala
    1370                 1375                 1380

Gln Asp  Gln Arg Ile Arg Trp  Tyr Leu Leu Ser Met  Gly Ser Asn
    1385                 1390                 1395

Glu Asn  Ile His Ser Ile His  Phe Ser Gly His Val  Phe Thr Val
    1400                 1405                 1410

Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
    1415                 1420                 1425

Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    1430                 1435                 1440

Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    1445                 1450                 1455

Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    1460                 1465                 1470

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
    1475                 1480                 1485

Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
    1490                 1495                 1500

Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
    1505                 1510                 1515

Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
    1520                 1525                 1530

Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    1535                 1540                 1545
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Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
    1550                1555                1560

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    1565                1570                1575

Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
    1580                1585                1590

Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1595                1600                1605

Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
    1610                1615                1620

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    1625                1630                1635

Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
    1640                1645                1650

Ser Lys  Ala Arg Leu His Leu  Gln Gly Arg Ser Asn  Ala Trp Arg
    1655                1660                1665

Pro Gln  Val Asn Asn Pro Lys  Glu Trp Leu Gln Val  Asp Phe Gln
    1670                1675                1680

Lys Thr  Met Lys Val Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser
    1685                1690                1695

Leu Leu  Thr Ser Met Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser
    1700                1705                1710

Gln Asp  Gly His Gln Trp Thr  Leu Phe Phe Gln Asn  Gly Lys Val
    1715                1720                1725

Lys Val  Phe Gln Gly Asn Gln  Asp Ser Phe Thr Pro  Val Val Asn
    1730                1735                1740

Ser Leu  Asp Pro Pro Leu Leu  Thr Arg Tyr Leu Arg  Ile His Pro
    1745                1750                1755

Gln Ser  Trp Val His Gln Ile  Ala Leu Arg Met Glu  Val Leu Gly
    1760                1765                1770

Cys Glu  Ala Gln Asp Leu Tyr  Gly Ala Pro Gly Thr  Ser Glu Ser
    1775                1780                1785

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
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Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
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Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Thr  Ser Thr Glu
    1835                 1840                 1845

Pro Ser  Glu Gly Ser Ala Pro  Gly Ser Pro Ala Gly  Ser Pro Thr
    1850                 1855                 1860

Ser Thr  Glu Glu Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1865                 1870                 1875

Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr
    1880                 1885                 1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Pro Ala Gly
    1895                 1900                 1905

Ser Pro  Thr Ser Thr Glu Glu  Gly Ser Pro Ala Gly  Ser Pro Thr
    1910                 1915                 1920

Ser Thr  Glu Glu Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    1925                 1930                 1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                 1945                 1950

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Thr  Ser Glu Ser
    1955                 1960                 1965

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1970                 1975                 1980

Ser Glu  Thr Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                 1990                 1995

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr
    2000                 2005                 2010

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Thr  Ser Thr Glu
    2015                 2020                 2025

Pro Ser  Glu Gly Ser Ala Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    2030                 2035                 2040

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2045                 2050                 2055
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Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Ala Ser
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Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala Thr
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Gly His His His His His His
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val

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Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
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Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
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Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
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405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Ala Pro Gly Ser Pro Ala Gly Ser
595 600 605

Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
610 615 620

Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Ala Ser
625 630 635 640

Ser Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met
645 650 655

His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys

660 665 670

Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr
675 680 685

Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met
690 695 700

Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val
705 710 715 720

Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn
725 730 735

Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser
740 745 750

Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile
755 760 765

Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe
770 775 780

Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
785 790 795 800

Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
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Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
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Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
885 890 895

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro
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915 920 925

Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro Pro Val Leu Lys Arg His

930 935 940

Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
945 950 955 960

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
965 970 975

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
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Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
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Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn
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Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe
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Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala
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Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr Lys Thr Tyr
1100 1105 1110

Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp Glu Phe
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Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu Lys
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Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val Cys His Thr
1145 1150 1155

Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr Val Gln Glu
1160 1165 1170

Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys Ser Trp Tyr
1175 1180 1185

Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Gly Ala Pro Thr Ser

1190 1195 1200

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Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
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Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
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Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
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Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
1310 1315 1320

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro
1325 1330 1335

Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Ala Pro Cys Asn Ile
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Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg Phe His Ala
1355 1360 1365

Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu Val Met Ala
1370 1375 1380

Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met Gly Ser Asn
1385 1390 1395

Glu Asn Ile His Ser Ile His Phe Ser Gly His Val Phe Thr Val
1400 1405 1410

Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn Leu Tyr Pro
1415 1420 1425

Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys Ala Gly Ile
1430 1435 1440

Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His Ala Gly Met

1445 1450 1455

Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln Thr Pro Leu
1460 1465 1470

Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile Thr Ala Ser
1475 1480 1485

Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg Leu His Tyr
1490 1495 1500

Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro Phe Ser Trp
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Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His Gly Ile Lys
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Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr Ile Ser Gln
1535 1540 1545

Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp Gln Thr Tyr
1550 1555 1560

Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe Gly Asn Val
1565 1570 1575

Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro Pro Ile Ile
1580 1585 1590

Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser Ile Arg Ser
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Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp Ala Gln Ile
1625 1630 1635

Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp Ser Pro
1640 1645 1650

Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg
1655 1660 1665

Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln
1670 1675 1680

Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser
1685 1690 1695

Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser

1700 1705 1710

Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val
1715 1720 1725

Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn
1730 1735 1740

Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro
1745 1750 1755

Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly
1760 1765 1770

Cys Glu Ala Gln Asp Leu Tyr Gly Ala Pro Gly Thr Ser Glu Ser
1775 1780 1785

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1835 1840 1845

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Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1865 1870 1875

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr
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1910 1915 1920

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1925 1930 1935

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1955 1960 1965

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1985 1990 1995

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr
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<211> 2095
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285
```

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Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560
```

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Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
            740                 745                 750

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
        755                 760                 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
    770                 775                 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785                 790                 795                 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                805                 810                 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            820                 825                 830
```

```
Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
        835                 840                 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    850                 855                 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865                 870                 875                 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095
```

```
Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1145                1150                1155

Pro Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    1160                1165                1170

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1175                1180                1185

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1190                1195                1200

Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1205                1210                1215

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1220                1225                1230

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Pro  Ala Gly Ser
    1235                1240                1245

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1250                1255                1260

Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1265                1270                1275

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1280                1285                1290

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ala Ser  Ser Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350
```

```
Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Gly  Ala Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    1475                1480                1485

Glu Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1490                1495                1500

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Ala Ser  Ser Ser Trp
    1505                1510                1515

Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
    1520                1525                1530

Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    1535                1540                1545

Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
    1550                1555                1560

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    1565                1570                1575

Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
    1580                1585                1590

Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1595                1600                1605
```

```
Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
    1610                1615                1620

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    1625                1630                1635

Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
    1640                1645                1650

Ser Lys  Ala Arg Leu His Leu  Gln Gly Arg Ser Asn  Ala Trp Arg
    1655                1660                1665

Pro Gln  Val Asn Asn Pro Lys  Glu Trp Leu Gln Val  Asp Phe Gln
    1670                1675                1680

Lys Thr  Met Lys Val Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser
    1685                1690                1695

Leu Leu  Thr Ser Met Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser
    1700                1705                1710

Gln Asp  Gly His Gln Trp Thr  Leu Phe Phe Gln Asn  Gly Lys Val
    1715                1720                1725

Lys Val  Phe Gln Gly Asn Gln  Asp Ser Phe Thr Pro  Val Val Asn
    1730                1735                1740

Ser Leu  Asp Pro Pro Leu Leu  Thr Arg Tyr Leu Arg  Ile His Pro
    1745                1750                1755

Gln Ser  Trp Val His Gln Ile  Ala Leu Arg Met Glu  Val Leu Gly
    1760                1765                1770

Cys Glu  Ala Gln Asp Leu Tyr  Gly Ala Pro Gly Thr  Ser Glu Ser
    1775                1780                1785

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1790                1795                1800

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1805                1810                1815

Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr
    1820                1825                1830

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Thr  Ser Thr Glu
    1835                1840                1845

Pro Ser  Glu Gly Ser Ala Pro  Gly Ser Pro Ala Gly  Ser Pro Thr
    1850                1855                1860
```

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Ser Thr  Glu Glu Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1865                1870                1875

Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr
    1880                1885                1890

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Pro Ala Gly
    1895                1900                1905

Ser Pro  Thr Ser Thr Glu Glu  Gly Ser Pro Ala Gly  Ser Pro Thr
    1910                1915                1920

Ser Thr  Glu Glu Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Thr  Ser Glu Ser
    1955                1960                1965

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1970                1975                1980

Ser Glu  Thr Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                1990                1995

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr
    2000                2005                2010

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Thr  Ser Thr Glu
    2015                2020                2025

Pro Ser  Glu Gly Ser Ala Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    2030                2035                2040

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2045                2050                2055

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Ala Ser
    2060                2065                2070

Ser Glu  Gln Lys Leu Ile Ser  Glu Glu Asp Leu Ser  Pro Ala Thr
    2075                2080                2085

Gly His  His His His His His
    2090                2095
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gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

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gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

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Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
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Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
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Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720
```

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Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
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Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
            740                 745                 750

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
        755                 760                 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
    770                 775                 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785                 790                 795                 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                805                 810                 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            820                 825                 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
        835                 840                 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    850                 855                 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865                 870                 875                 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990
```

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Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
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Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
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Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1145                1150                1155

Pro Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    1160                1165                1170

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1175                1180                1185

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1190                1195                1200

Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1205                1210                1215

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
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Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Pro  Ala Gly Ser
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Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
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Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1265                 1270                 1275

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1280                 1285                 1290

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ala Ser  Ser Ala Pro
    1295                 1300                 1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                 1315                 1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                 1330                 1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                 1345                 1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                 1360                 1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                 1375                 1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                 1390                 1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                 1405                 1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                 1420                 1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                 1435                 1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                 1450                 1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                 1465                 1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                 1480                 1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                 1495                 1500
```

```
Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Gly
    1565                1570                1575

Ala Pro  Gly Ser Pro Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly
    1580                1585                1590

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro
    1595                1600                1605

Ala Thr  Ser Gly Ser Glu Thr  Pro Ala Ser Ser Asn  Ser Cys Ser
    1610                1615                1620

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    1625                1630                1635

Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
    1640                1645                1650

Ser Lys  Ala Arg Leu His Leu  Gln Gly Arg Ser Asn  Ala Trp Arg
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Pro Gln  Val Asn Asn Pro Lys  Glu Trp Leu Gln Val  Asp Phe Gln
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Lys Thr  Met Lys Val Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser
    1685                1690                1695

Leu Leu  Thr Ser Met Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser
    1700                1705                1710

Gln Asp  Gly His Gln Trp Thr  Leu Phe Phe Gln Asn  Gly Lys Val
    1715                1720                1725

Lys Val  Phe Gln Gly Asn Gln  Asp Ser Phe Thr Pro  Val Val Asn
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Ser Leu  Asp Pro Pro Leu Leu  Thr Arg Tyr Leu Arg  Ile His Pro
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```

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Gln Ser  Trp Val His Gln Ile  Ala Leu Arg Met Glu  Val Leu Gly
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Cys Glu  Ala Gln Asp Leu Tyr  Gly Ala Pro Gly Thr  Ser Glu Ser
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Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1790                1795                1800

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1805                1810                1815

Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr
    1820                1825                1830

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Thr  Ser Thr Glu
    1835                1840                1845

Pro Ser  Glu Gly Ser Ala Pro  Gly Ser Pro Ala Gly  Ser Pro Thr
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Ser Thr  Glu Glu Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
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Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Pro Ala Gly
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Ser Pro  Thr Ser Thr Glu Glu  Gly Ser Pro Ala Gly  Ser Pro Thr
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Ser Thr  Glu Glu Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    1925                1930                1935

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1940                1945                1950

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Thr  Ser Glu Ser
    1955                1960                1965

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1970                1975                1980

Ser Glu  Thr Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1985                1990                1995

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr
    2000                2005                2010
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Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
2015 2020 2025

Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
2030 2035 2040

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2045 2050 2055

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

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Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335
```

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Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605
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Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
            740                 745                 750

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
        755                 760                 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
    770                 775                 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785                 790                 795                 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                805                 810                 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            820                 825                 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
        835                 840                 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    850                 855                 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865                 870                 875                 880
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Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140
```

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Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
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Pro Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    1160                1165                1170

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1175                1180                1185

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1190                1195                1200

Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1205                1210                1215

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1220                1225                1230

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Pro  Ala Gly Ser
    1235                1240                1245

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1250                1255                1260

Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1265                1270                1275

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1280                1285                1290

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ala Ser  Ser Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395
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Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
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Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Gly Ala  Pro Gly Ser
    1625                1630                1635

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
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```

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Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
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Ser Glu  Thr Pro Ala Ser Ser  Glu Trp Leu Gln Val  Asp Phe Gln
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Lys Thr  Met Lys Val Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser
    1685                 1690                 1695

Leu Leu  Thr Ser Met Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser
    1700                 1705                 1710

Gln Asp  Gly His Gln Trp Thr  Leu Phe Phe Gln Asn  Gly Lys Val
    1715                 1720                 1725

Lys Val  Phe Gln Gly Asn Gln  Asp Ser Phe Thr Pro  Val Val Asn
    1730                 1735                 1740

Ser Leu  Asp Pro Pro Leu Leu  Thr Arg Tyr Leu Arg  Ile His Pro
    1745                 1750                 1755

Gln Ser  Trp Val His Gln Ile  Ala Leu Arg Met Glu  Val Leu Gly
    1760                 1765                 1770

Cys Glu  Ala Gln Asp Leu Tyr  Gly Ala Pro Gly Thr  Ser Glu Ser
    1775                 1780                 1785

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1790                 1795                 1800

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1805                 1810                 1815

Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr
    1820                 1825                 1830

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Thr  Ser Thr Glu
    1835                 1840                 1845

Pro Ser  Glu Gly Ser Ala Pro  Gly Ser Pro Ala Gly  Ser Pro Thr
    1850                 1855                 1860

Ser Thr  Glu Glu Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1865                 1870                 1875

Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Pro Ala Gly
    1895                 1900                 1905
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Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1940 1945 1950

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser
1955 1960 1965

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
1970 1975 1980

Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1985 1990 1995

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr
2000 2005 2010

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
2015 2020 2025

Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
2030 2035 2040

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
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Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Ala Ser
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
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225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
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Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys

500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
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Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
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Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
740 745 750

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
755 760 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly

770 775 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785 790 795 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
805 810 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
820 825 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
835 840 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
850 855 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865 870 875 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
885 890 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
900 905 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
915 920 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
930 935 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945 950 955 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
965 970 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
980 985 990

Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
995 1000 1005

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
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Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
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Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg

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Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
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Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
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Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
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Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1145                1150                1155

Pro Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
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Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1175                1180                1185

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
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Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
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Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
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Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Pro  Ala Gly Ser
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Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
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Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
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Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
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Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ala Ser  Ser Ala Pro
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Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
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Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
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Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
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Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
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Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
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Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
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Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
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Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
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Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
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Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
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Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
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Gly Ala  Pro Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
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Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
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Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Ala Ser Ser  Gly Lys Val
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Lys Val  Phe Gln Gly Asn Gln  Asp Ser Phe Thr Pro  Val Val Asn
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Ser Leu  Asp Pro Pro Leu Leu  Thr Arg Tyr Leu Arg  Ile His Pro
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Gln Ser  Trp Val His Gln Ile  Ala Leu Arg Met Glu  Val Leu Gly
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Cys Glu  Ala Gln Asp Leu Tyr  Gly Ala Pro Gly Thr  Ser Glu Ser
    1775                1780                1785

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
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Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
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Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
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Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1865 1870 1875

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr
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Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
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Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
1925 1930 1935

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1940 1945 1950

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser
1955 1960 1965

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
1970 1975 1980

Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1985 1990 1995

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr
2000 2005 2010

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
2015 2020 2025

Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
2030 2035 2040

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
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Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Ala Ser

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<212> PRT
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Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
35 40 45

Pro Ala Thr Ser Gly Ser Glu Thr Pro Ala Ser Ser Ser Asp Leu Gly
50 55 60

Glu Leu Pro Val Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe
65 70 75 80

Pro Phe Asn Thr Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe
85 90 95

Thr Asp His Leu Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly
100 105 110

Leu Leu Gly Pro Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile
115 120 125

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Thr Leu Lys Asn Met Ala Ser His Pro Val Ser Leu His Ala Val Gly
    130                 135                 140

Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr
145                 150                 155                 160

Ser Gln Arg Glu Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His
                165                 170                 175

Thr Tyr Val Trp Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp
            180                 185                 190

Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys
        195                 200                 205

Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly
    210                 215                 220

Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu
225                 230                 235                 240

Phe Ala Val Phe Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn
                245                 250                 255

Ser Leu Met Gln Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys
            260                 265                 270

Met His Thr Val Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile
        275                 280                 285

Gly Cys His Arg Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr
    290                 295                 300

Thr Pro Glu Val His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val
305                 310                 315                 320

Arg Asn His Arg Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu
                325                 330                 335

Thr Ala Gln Thr Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys
            340                 345                 350

His Ile Ser Ser His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val
        355                 360                 365

Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu
    370                 375                 380

Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val
385                 390                 395                 400
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Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val
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Ala Lys Lys His Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu
            420                 425                 430

Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser
        435                 440                 445

Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys
    450                 455                 460

Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr
465                 470                 475                 480

Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr
                485                 490                 495

Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser
            500                 505                 510

Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu
        515                 520                 525

Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro
    530                 535                 540

Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu
545                 550                 555                 560

Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser
                565                 570                 575

Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro
            580                 585                 590

Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile
        595                 600                 605

Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn
    610                 615                 620

Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro
625                 630                 635                 640

Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met
                645                 650                 655

His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys
            660                 665                 670
```

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Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr
        675                 680                 685

Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met
    690                 695                 700

Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val
705                 710                 715                 720

Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn
                725                 730                 735

Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser
            740                 745                 750

Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile
        755                 760                 765

Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe
    770                 775                 780

Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
785                 790                 795                 800

Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
                805                 810                 815

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
            820                 825                 830

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
        835                 840                 845

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
    850                 855                 860

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
865                 870                 875                 880

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
                885                 890                 895

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro
            900                 905                 910

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
        915                 920                 925

Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro Pro Val Leu Lys Arg His
    930                 935                 940
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Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
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Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
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Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
            980                 985                 990

Thr Arg His Tyr Phe Ile Ala Ala  Val Glu Arg Leu Trp  Asp Tyr Gly
        995                 1000                1005

Met Ser  Ser Ser Pro His Val  Leu Arg Asn Arg Ala  Gln Ser Gly
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Ser Val  Pro Gln Phe Lys Lys  Val Val Phe Gln Glu  Phe Thr Asp
    1025                1030                1035

Gly Ser  Phe Thr Gln Pro Leu  Tyr Arg Gly Glu Leu  Asn Glu His
    1040                1045                1050

Leu Gly  Leu Leu Gly Pro Tyr  Ile Arg Ala Glu Val  Glu Asp Asn
    1055                1060                1065

Ile Met  Val Thr Phe Arg Asn  Gln Ala Ser Arg Pro  Tyr Ser Phe
    1070                1075                1080

Tyr Ser  Ser Leu Ile Ser Tyr  Glu Glu Asp Gln Arg  Gln Gly Ala
    1085                1090                1095

Glu Pro  Arg Lys Asn Phe Val  Lys Pro Asn Glu Thr  Lys Thr Tyr
    1100                1105                1110

Phe Trp  Lys Val Gln His His  Met Ala Pro Thr Lys  Asp Glu Phe
    1115                1120                1125

Asp Cys  Lys Ala Trp Ala Tyr  Phe Ser Asp Val Asp  Leu Glu Lys
    1130                1135                1140

Asp Val  His Ser Gly Leu Ile  Gly Pro Leu Leu Val  Cys His Thr
    1145                1150                1155

Asn Thr  Leu Asn Pro Ala His  Gly Arg Gln Val Thr  Val Gln Glu
    1160                1165                1170

Phe Ala  Leu Phe Phe Thr Ile  Phe Asp Glu Thr Lys  Ser Trp Tyr
    1175                1180                1185

Phe Thr  Glu Asn Met Glu Arg  Asn Cys Arg Ala Pro  Cys Asn Ile
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Gln Met  Glu Asp Pro Thr Phe  Lys Glu Asn Tyr Arg  Phe His Ala
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Ile Asn  Gly Tyr Ile Met Asp  Thr Leu Pro Gly Leu  Val Met Ala
    1220                 1225                 1230

Gln Asp  Gln Arg Ile Arg Trp  Tyr Leu Leu Ser Met  Gly Ser Asn
    1235                 1240                 1245

Glu Asn  Ile His Ser Ile His  Phe Ser Gly His Val  Phe Thr Val
    1250                 1255                 1260

Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
    1265                 1270                 1275

Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    1280                 1285                 1290

Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    1295                 1300                 1305

Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    1310                 1315                 1320

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
    1325                 1330                 1335

Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
    1340                 1345                 1350

Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
    1355                 1360                 1365

Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
    1370                 1375                 1380

Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    1385                 1390                 1395

Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
    1400                 1405                 1410

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    1415                 1420                 1425

Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
    1430                 1435                 1440

Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1445                 1450                 1455
```

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Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
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Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    1475                1480                1485

Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
    1490                1495                1500

Ser Lys  Ala Arg Leu His Leu  Gln Gly Arg Ser Asn  Ala Trp Arg
    1505                1510                1515

Pro Gln  Val Asn Asn Pro Lys  Glu Trp Leu Gln Val  Asp Phe Gln
    1520                1525                1530

Lys Thr  Met Lys Val Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser
    1535                1540                1545

Leu Leu  Thr Ser Met Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser
    1550                1555                1560

Gln Asp  Gly His Gln Trp Thr  Leu Phe Phe Gln Asn  Gly Lys Val
    1565                1570                1575

Lys Val  Phe Gln Gly Asn Gln  Asp Ser Phe Thr Pro  Val Val Asn
    1580                1585                1590

Ser Leu  Asp Pro Pro Leu Leu  Thr Arg Tyr Leu Arg  Ile His Pro
    1595                1600                1605

Gln Ser  Trp Val His Gln Ile  Ala Leu Arg Met Glu  Val Leu Gly
    1610                1615                1620

Cys Glu  Ala Gln Asp Leu Tyr  Gly Ala Pro Gly Thr  Ser Glu Ser
    1625                1630                1635

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1640                1645                1650

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1655                1660                1665

Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Thr  Ser Thr Glu
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Pro Ser  Glu Gly Ser Ala Pro  Gly Ser Pro Ala Gly  Ser Pro Thr
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Ser Thr  Glu Glu Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
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Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Pro Ala Gly
    1745                1750                1755

Ser Pro  Thr Ser Thr Glu Glu  Gly Ser Pro Ala Gly  Ser Pro Thr
    1760                1765                1770

Ser Thr  Glu Glu Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    1775                1780                1785

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1790                1795                1800

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Thr  Ser Glu Ser
    1805                1810                1815

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1820                1825                1830

Ser Glu  Thr Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1835                1840                1845

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr
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Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Thr  Ser Thr Glu
    1865                1870                1875

Pro Ser  Glu Gly Ser Ala Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1880                1885                1890

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1895                1900                1905

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Ala Ser
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Ser Glu  Gln Lys Leu Ile Ser  Glu Glu Asp Leu Ser  Pro Ala Thr
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Gly His  His His His His His
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tttcagggaa atcaagactc cttcacacct gtggtgaact ctctagaccc accgttactg 4800

actcgctacc ttcgaattca cccccagagt tgggtgcacc agattgccct gaggatggag 4860

gttctgggct gcgaggcaca ggacctctac ggcgcgccag gtacctcaga gtctgctacc 4920

cccgagtcag ggccaggatc agagccagcc acctccgggt ctgagacacc cgggacttcc 4980

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gagagtgcca cccccgagag tggcccaggc actagtaccg agccctctga aggcagtgcg 5760

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<210> 1373
<211> 1945
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg

225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Gly Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr
405 410 415

Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
420 425 430

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Ala Ser Ser Asp Arg Ser
435 440 445

Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys
450 455 460

Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr
465 470 475 480

Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr
485 490 495

Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser

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            500                 505                 510

Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu
        515                 520                 525

Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro
    530                 535                 540

Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu
545                 550                 555                 560

Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser
                565                 570                 575

Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro
            580                 585                 590

Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile
        595                 600                 605

Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn
    610                 615                 620

Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro
625                 630                 635                 640

Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met
                645                 650                 655

His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys
            660                 665                 670

Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr
        675                 680                 685

Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met
    690                 695                 700

Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val
705                 710                 715                 720

Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn
                725                 730                 735

Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser
            740                 745                 750

Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile
        755                 760                 765

Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe
```

770 775 780

Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
785 790 795 800

Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
805 810 815

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
820 825 830

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
835 840 845

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
850 855 860

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
865 870 875 880

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
885 890 895

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro
900 905 910

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
915 920 925

Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro Pro Val Leu Lys Arg His
930 935 940

Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
945 950 955 960

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
965 970 975

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
980 985 990

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
995 1000 1005

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly
1010 1015 1020

Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp
1025 1030 1035

Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His

1040 1045 1050

Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn
1055 1060 1065

Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe
1070 1075 1080

Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala
1085 1090 1095

Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr Lys Thr Tyr
1100 1105 1110

Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp Glu Phe
1115 1120 1125

Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu Lys
1130 1135 1140

Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val Cys His Thr
1145 1150 1155

Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr Val Gln Glu
1160 1165 1170

Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys Ser Trp Tyr
1175 1180 1185

Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro Cys Asn Ile
1190 1195 1200

Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg Phe His Ala
1205 1210 1215

Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu Val Met Ala
1220 1225 1230

Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met Gly Ser Asn
1235 1240 1245

Glu Asn Ile His Ser Ile His Phe Ser Gly His Val Phe Thr Val
1250 1255 1260

Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn Leu Tyr Pro
1265 1270 1275

Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys Ala Gly Ile
1280 1285 1290

Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His Ala Gly Met

1295 1300 1305

Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln Thr Pro Leu
1310 1315 1320

Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile Thr Ala Ser
1325 1330 1335

Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg Leu His Tyr
1340 1345 1350

Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro Phe Ser Trp
1355 1360 1365

Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His Gly Ile Lys
1370 1375 1380

Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr Ile Ser Gln
1385 1390 1395

Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp Gln Thr Tyr
1400 1405 1410

Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe Gly Asn Val
1415 1420 1425

Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro Pro Ile Ile
1430 1435 1440

Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser Ile Arg Ser
1445 1450 1455

Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn Ser Cys Ser
1460 1465 1470

Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp Ala Gln Ile
1475 1480 1485

Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp Ser Pro
1490 1495 1500

Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg
1505 1510 1515

Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln
1520 1525 1530

Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser
1535 1540 1545

Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser

1550 1555 1560

Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val
1565 1570 1575

Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn
1580 1585 1590

Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro
1595 1600 1605

Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly
1610 1615 1620

Cys Glu Ala Gln Asp Leu Tyr Gly Ala Pro Gly Thr Ser Glu Ser
1625 1630 1635

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
1640 1645 1650

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1655 1660 1665

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr
1670 1675 1680

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
1685 1690 1695

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
1700 1705 1710

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1715 1720 1725

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr
1730 1735 1740

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly
1745 1750 1755

Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
1760 1765 1770

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
1775 1780 1785

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1790 1795 1800

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser

1805 1810 1815

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
1820 1825 1830

Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1835 1840 1845

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr
1850 1855 1860

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
1865 1870 1875

Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly
1880 1885 1890

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1895 1900 1905

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Ala Ser
1910 1915 1920

Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala Thr
1925 1930 1935

Gly His His His His His His
1940 1945

<210> 1374
<211> 5838
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

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atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

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acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

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gagagtgcca cccccgagag tggcccaggc actagtaccg agccctctga aggcagtgcg 5760

ccagcctcga gcgaacaaaa actcatctca gaagaggatc tgtcacctgc aaccggtcat 5820

catcaccatc accattga 5838

<210> 1375
<211> 1948
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

```
Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335
```

```
Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605
```

```
Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
            740                 745                 750

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
        755                 760                 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
    770                 775                 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785                 790                 795                 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                805                 810                 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            820                 825                 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
        835                 840                 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    850                 855                 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865                 870                 875                 880
```

```
Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Gly Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr
            980                 985                 990

Glu Glu Gly Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
        995                 1000                1005

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Ala Ser  Ser Arg Ala
    1010                1015                1020

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1025                1030                1035

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1040                1045                1050

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1055                1060                1065

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1070                1075                1080

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1085                1090                1095

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1100                1105                1110

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1115                1120                1125

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1130                1135                1140
```

```
Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1145                1150                1155

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1160                1165                1170

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1175                1180                1185

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1190                1195                1200

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1205                1210                1215

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1220                1225                1230

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1235                1240                1245

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1250                1255                1260

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1265                1270                1275

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1280                1285                1290

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1295                1300                1305

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1310                1315                1320

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1325                1330                1335

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1340                1345                1350

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1355                1360                1365

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1370                1375                1380

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1385                1390                1395
```

```
Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1400                1405                1410

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1415                1420                1425

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1430                1435                1440

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1445                1450                1455

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1460                1465                1470

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1475                1480                1485

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1490                1495                1500

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1505                1510                1515

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1520                1525                1530

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1535                1540                1545

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1550                1555                1560

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1565                1570                1575

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1580                1585                1590

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1595                1600                1605

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1610                1615                1620

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1625                1630                1635

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1640                1645                1650
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```
Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1655                 1660                 1665

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1670                 1675                 1680

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1685                 1690                 1695

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1700                 1705                 1710

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1715                 1720                 1725

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1730                 1735                 1740

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1745                 1750                 1755

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    1760                 1765                 1770

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    1775                 1780                 1785

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1790                 1795                 1800

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1805                 1810                 1815

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1820                 1825                 1830

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1835                 1840                 1845

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    1850                 1855                 1860

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    1865                 1870                 1875

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Glu Pro Ala
    1880                 1885                 1890

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1895                 1900                 1905
```

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
1910 1915 1920

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Ala Thr Gly His His His His His His
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<211> 5847
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atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

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Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445
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Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720
```

```
Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
            740                 745                 750

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
        755                 760                 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
    770                 775                 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785                 790                 795                 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                805                 810                 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            820                 825                 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
        835                 840                 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    850                 855                 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865                 870                 875                 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990
```

```
Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1145                1150                1155

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr
    1160                1165                1170

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1175                1180                1185

Thr Ser  Gly Ser Glu Thr Pro  Ala Ser Ser Ala Pro  Cys Asn Ile
    1190                1195                1200

Gln Met  Glu Asp Pro Thr Phe  Lys Glu Asn Tyr Arg  Phe His Ala
    1205                1210                1215

Ile Asn  Gly Tyr Ile Met Asp  Thr Leu Pro Gly Leu  Val Met Ala
    1220                1225                1230

Gln Asp  Gln Arg Ile Arg Trp  Tyr Leu Leu Ser Met  Gly Ser Asn
    1235                1240                1245
```

```
Glu Asn  Ile His Ser Ile His  Phe Ser Gly His Val  Phe Thr Val
    1250                1255                1260

Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
    1265                1270                1275

Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    1280                1285                1290

Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    1295                1300                1305

Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    1310                1315                1320

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
    1325                1330                1335

Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
    1340                1345                1350

Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
    1355                1360                1365

Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
    1370                1375                1380

Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    1385                1390                1395

Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
    1400                1405                1410

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    1415                1420                1425

Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
    1430                1435                1440

Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1445                1450                1455

Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
    1460                1465                1470

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    1475                1480                1485

Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
    1490                1495                1500
```

```
Ser Lys  Ala Arg Leu His Leu  Gln Gly Arg Ser Asn  Ala Trp Arg
    1505                 1510                 1515

Pro Gln  Val Asn Asn Pro Lys  Glu Trp Leu Gln Val  Asp Phe Gln
    1520                 1525                 1530

Lys Thr  Met Lys Val Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser
    1535                 1540                 1545

Leu Leu  Thr Ser Met Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser
    1550                 1555                 1560

Gln Asp  Gly His Gln Trp Thr  Leu Phe Phe Gln Asn  Gly Lys Val
    1565                 1570                 1575

Lys Val  Phe Gln Gly Asn Gln  Asp Ser Phe Thr Pro  Val Val Asn
    1580                 1585                 1590

Ser Leu  Asp Pro Pro Leu Leu  Thr Arg Tyr Leu Arg  Ile His Pro
    1595                 1600                 1605

Gln Ser  Trp Val His Gln Ile  Ala Leu Arg Met Glu  Val Leu Gly
    1610                 1615                 1620

Cys Glu  Ala Gln Asp Leu Tyr  Gly Ala Pro Gly Thr  Ser Glu Ser
    1625                 1630                 1635

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1640                 1645                 1650

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1655                 1660                 1665

Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr
    1670                 1675                 1680

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Thr  Ser Thr Glu
    1685                 1690                 1695

Pro Ser  Glu Gly Ser Ala Pro  Gly Ser Pro Ala Gly  Ser Pro Thr
    1700                 1705                 1710

Ser Thr  Glu Glu Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1715                 1720                 1725

Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr
    1730                 1735                 1740

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Pro Ala Gly
    1745                 1750                 1755
```

```
Ser Pro  Thr Ser Thr Glu Glu  Gly Ser Pro Ala Gly  Ser Pro Thr
    1760                1765                1770

Ser Thr  Glu Glu Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    1775                1780                1785

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1790                1795                1800

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Thr  Ser Glu Ser
    1805                1810                1815

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1820                1825                1830

Ser Glu  Thr Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1835                1840                1845

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr
    1850                1855                1860

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Thr  Ser Thr Glu
    1865                1870                1875

Pro Ser  Glu Gly Ser Ala Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1880                1885                1890

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1895                1900                1905

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Ala Ser
    1910                1915                1920

Ser Glu  Gln Lys Leu Ile Ser  Glu Glu Asp Leu Ser  Pro Ala Thr
    1925                1930                1935

Gly His  His His His His His
    1940                1945
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<211> 5838
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atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

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<211> 2053
<212> PRT
<213> Artificial Sequence

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285
```

```
Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
                405                 410                 415

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            420                 425                 430

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
        435                 440                 445

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    450                 455                 460

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
465                 470                 475                 480

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
                485                 490                 495

Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            500                 505                 510

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
        515                 520                 525

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
    530                 535                 540

Pro Glu Ser Gly Pro Gly Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560
```

```
Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830
```

```
Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
        835                 840                 845

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
                885                 890                 895

Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            900                 905                 910

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        915                 920                 925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    930                 935                 940

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
945                 950                 955                 960

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                965                 970                 975

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            980                 985                 990

Pro Gly Thr Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
        995                 1000                1005

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1010                1015                1020

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1025                1030                1035

Ser Ala  Pro Gly Ala Ser Ser  Pro Pro Val Leu Lys  Arg His Gln
    1040                1045                1050

Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080

Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
    1085                1090                1095
```

```
Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
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Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                1120                1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                1135                1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                1150                1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                1165                1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                1180                1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                1195                1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                1210                1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                1225                1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                1240                1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                1255                1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                1270                1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                1285                1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350
```

```
Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                 1360                 1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                 1375                 1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                 1390                 1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                 1405                 1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                 1420                 1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                 1435                 1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                 1450                 1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                 1465                 1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                 1480                 1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                 1495                 1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                 1510                 1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                 1525                 1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                 1540                 1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                 1555                 1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                 1570                 1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                 1585                 1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                 1600                 1605
```

```
Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1730                1735                1740

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1745                1750                1755

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1760                1765                1770

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1775                1780                1785

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1790                1795                1800

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1805                1810                1815

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1820                1825                1830

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1835                1840                1845

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1850                1855                1860
```

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
1865 1870 1875

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
1880 1885 1890

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1895 1900 1905

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1910 1915 1920

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1925 1930 1935

Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
1940 1945 1950

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
1955 1960 1965

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
1970 1975 1980

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
1985 1990 1995

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
2000 2005 2010

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2015 2020 2025

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Ala Thr Gly His His His His His His
2045 2050

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<400> 1381

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            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240
```

```
Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510
```

```
Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
            740                 745                 750

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
        755                 760                 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
    770                 775                 780
```

```
Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785                 790                 795                 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                805                 810                 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            820                 825                 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
        835                 840                 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    850                 855                 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865                 870                 875                 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Gly Ala Pro Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
            980                 985                 990

Pro Gly Ser Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser
        995                 1000                1005

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
    1010                1015                1020

Ser Gly  Ser Glu Thr Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1025                1030                1035

Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1040                1045                1050
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Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Pro
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Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Ser Pro  Ala Gly Ser
    1070                 1075                 1080

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1085                 1090                 1095

Thr Glu  Glu Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1100                 1105                 1110

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Ala Ser
    1115                 1120                 1125

Ser Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    1130                 1135                 1140

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    1145                 1150                 1155

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1160                 1165                 1170

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1175                 1180                 1185

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1190                 1195                 1200

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1205                 1210                 1215

Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1220                 1225                 1230

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1235                 1240                 1245

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1250                 1255                 1260

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1265                 1270                 1275

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1280                 1285                 1290

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1295                 1300                 1305
```

```
Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1310                 1315                 1320

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1325                 1330                 1335

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1340                 1345                 1350

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1355                 1360                 1365

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1370                 1375                 1380

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1385                 1390                 1395

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1400                 1405                 1410

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1415                 1420                 1425

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1430                 1435                 1440

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1445                 1450                 1455

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1460                 1465                 1470

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1475                 1480                 1485

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1490                 1495                 1500

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1505                 1510                 1515

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1520                 1525                 1530

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1535                 1540                 1545

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1550                 1555                 1560
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His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1565                1570                1575

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1580                1585                1590

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1595                1600                1605

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1610                1615                1620

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1625                1630                1635

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1640                1645                1650

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1655                1660                1665

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1670                1675                1680

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1685                1690                1695

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    1700                1705                1710

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    1715                1720                1725

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    1730                1735                1740

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1745                1750                1755

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    1760                1765                1770

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1775                1780                1785

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1790                1795                1800

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    1805                1810                1815
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Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    1820                 1825                 1830

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1835                 1840                 1845

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1850                 1855                 1860

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser
    1865                 1870                 1875

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Thr Glu
    1880                 1885                 1890

Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1895                 1900                 1905

Glu Ser  Gly Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1910                 1915                 1920

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1925                 1930                 1935

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Ser  Glu Pro Ala
    1940                 1945                 1950

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Pro Ala Gly  Ser Pro Thr
    1955                 1960                 1965

Ser Thr  Glu Glu Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    1970                 1975                 1980

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    1985                 1990                 1995

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    2000                 2005                 2010

Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu
    2015                 2020                 2025

Gly Ser  Ala Pro Ala Ser Ser  Glu Gln Lys Leu Ile  Ser Glu Glu
    2030                 2035                 2040

Asp Leu  Ser Pro Ala Thr Gly  His His His His His  His
    2045                 2050                 2055
```

<210> 1382
<211> 6171
<212> DNA

<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

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atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

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catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

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gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

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<211> 2053
<212> PRT
<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic polypeptide

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20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

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His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460
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Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735
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Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
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Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
        755                 760                 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
    770                 775                 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785                 790                 795                 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                805                 810                 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            820                 825                 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
        835                 840                 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    850                 855                 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865                 870                 875                 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005
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```
Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
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Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1145                1150                1155

Pro Ser  Pro Ala Gly Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser
    1160                1165                1170

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro
    1175                1180                1185

Ser Glu  Gly Ser Ala Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1190                1195                1200

Thr Glu  Glu Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1205                1210                1215

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1220                1225                1230

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
    1235                1240                1245

Ser Gly  Ser Glu Thr Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1250                1255                1260
```

```
Glu Thr  Pro Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1265                1270                1275

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1280                1285                1290

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ala Ser  Ser Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515
```

```
Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
    1730                1735                1740

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1745                1750                1755

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1760                1765                1770
```

```
Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1775                1780                1785

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1790                1795                1800

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1805                1810                1815

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1820                1825                1830

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1835                1840                1845

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1850                1855                1860

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    1865                1870                1875

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    1880                1885                1890

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1895                1900                1905

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1910                1915                1920

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1925                1930                1935

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1940                1945                1950

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    1955                1960                1965

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    1970                1975                1980

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Glu Pro Ala
    1985                1990                1995

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2000                2005                2010

Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    2015                2020                2025
```

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
2030 2035 2040

Pro Ala Thr Gly His His His His His His
2045 2050

<210> 1384
<211> 6162
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

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ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

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gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

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gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

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gatctgtcac ctgcaaccgg tcatcatcac catcaccatt ga 6162

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<211> 2197
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
20 25 30

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
35 40 45

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly
50 55 60

Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
65 70 75 80

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
85 90 95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
100 105 110

Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
115 120 125

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
130 135 140

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser

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145                 150                 155                 160

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
                165                 170                 175

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro
            180                 185                 190

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
        195                 200                 205

Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
    210                 215                 220

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
225                 230                 235                 240

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
                245                 250                 255

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
            260                 265                 270

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
        275                 280                 285

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
    290                 295                 300

Ala Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
305                 310                 315                 320

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
                325                 330                 335

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
            340                 345                 350

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
        355                 360                 365

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
    370                 375                 380

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
385                 390                 395                 400

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
                405                 410                 415

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
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420 425 430

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
435 440 445

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
450 455 460

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
465 470 475 480

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
485 490 495

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
500 505 510

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
515 520 525

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
530 535 540

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
545 550 555 560

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
565 570 575

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
580 585 590

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
595 600 605

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
610 615 620

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
625 630 635 640

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
645 650 655

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
660 665 670

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
675 680 685

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln

690 695 700

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
705 710 715 720

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
725 730 735

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
740 745 750

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
755 760 765

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
770 775 780

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
785 790 795 800

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
805 810 815

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
820 825 830

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
835 840 845

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
850 855 860

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
865 870 875 880

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
885 890 895

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
900 905 910

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
915 920 925

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
930 935 940

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
945 950 955 960

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met

965 970 975

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
980 985 990

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
995 1000 1005

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr
1010 1015 1020

Leu Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln
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Asn Gly Ala Pro Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
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Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
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Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala
1070 1075 1080

Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
1085 1090 1095

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
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Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
1115 1120 1125

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro
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Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
1145 1150 1155

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
1160 1165 1170

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser
1175 1180 1185

Ser Pro Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr
1190 1195 1200

Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
1205 1210 1215

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp

1220 1225 1230

Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr
1235 1240 1245

Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser
1250 1255 1260

Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro
1265 1270 1275

Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe
1280 1285 1290

Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
1295 1300 1305

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val
1310 1315 1320

Thr Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser
1325 1330 1335

Leu Ile Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg
1340 1345 1350

Lys Asn Phe Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys
1355 1360 1365

Val Gln His His Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys
1370 1375 1380

Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu Lys Asp Val His
1385 1390 1395

Ser Gly Leu Ile Gly Pro Leu Leu Val Cys His Thr Asn Thr Leu
1400 1405 1410

Asn Pro Ala His Gly Arg Gln Val Thr Val Gln Glu Phe Ala Leu
1415 1420 1425

Phe Phe Thr Ile Phe Asp Glu Thr Lys Ser Trp Tyr Phe Thr Glu
1430 1435 1440

Asn Met Glu Arg Asn Cys Arg Ala Pro Cys Asn Ile Gln Met Glu
1445 1450 1455

Asp Pro Thr Phe Lys Glu Asn Tyr Arg Phe His Ala Ile Asn Gly
1460 1465 1470

Tyr Ile Met Asp Thr Leu Pro Gly Leu Val Met Ala Gln Asp Gln

1475 1480 1485

Arg Ile Arg Trp Tyr Leu Leu Ser Met Gly Ser Asn Glu Asn Ile
1490 1495 1500

His Ser Ile His Phe Ser Gly His Val Phe Thr Val Arg Lys Lys
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Glu Glu Tyr Lys Met Ala Leu Tyr Asn Leu Tyr Pro Gly Val Phe
1520 1525 1530

Glu Thr Val Glu Met Leu Pro Ser Lys Ala Gly Ile Trp Arg Val
1535 1540 1545

Glu Cys Leu Ile Gly Glu His Leu His Ala Gly Met Ser Thr Leu
1550 1555 1560

Phe Leu Val Tyr Ser Asn Lys Cys Gln Thr Pro Leu Gly Met Ala
1565 1570 1575

Ser Gly His Ile Arg Asp Phe Gln Ile Thr Ala Ser Gly Gln Tyr
1580 1585 1590

Gly Gln Trp Ala Pro Lys Leu Ala Arg Leu His Tyr Ser Gly Ser
1595 1600 1605

Ile Asn Ala Trp Ser Thr Lys Glu Pro Phe Ser Trp Ile Lys Val
1610 1615 1620

Asp Leu Leu Ala Pro Met Ile Ile His Gly Ile Lys Thr Gln Gly
1625 1630 1635

Ala Arg Gln Lys Phe Ser Ser Leu Tyr Ile Ser Gln Phe Ile Ile
1640 1645 1650

Met Tyr Ser Leu Asp Gly Lys Lys Trp Gln Thr Tyr Arg Gly Asn
1655 1660 1665

Ser Thr Gly Thr Leu Met Val Phe Phe Gly Asn Val Asp Ser Ser
1670 1675 1680

Gly Ile Lys His Asn Ile Phe Asn Pro Pro Ile Ile Ala Arg Tyr
1685 1690 1695

Ile Arg Leu His Pro Thr His Tyr Ser Ile Arg Ser Thr Leu Arg
1700 1705 1710

Met Glu Leu Met Gly Cys Asp Leu Asn Ser Cys Ser Met Pro Leu
1715 1720 1725

Gly Met Glu Ser Lys Ala Ile Ser Asp Ala Gln Ile Thr Ala Ser

1730 1735 1740

Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp Ser Pro Ser Lys Ala
1745 1750 1755

Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg Pro Gln Val
1760 1765 1770

Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln Lys Thr Met
1775 1780 1785

Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser Leu Leu Thr
1790 1795 1800

Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly
1805 1810 1815

His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val Lys Val Phe
1820 1825 1830

Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn Ser Leu Asp
1835 1840 1845

Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro Gln Ser Trp
1850 1855 1860

Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly Cys Glu Ala
1865 1870 1875

Gln Asp Leu Tyr Gly Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
1880 1885 1890

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1895 1900 1905

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
1910 1915 1920

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
1925 1930 1935

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
1940 1945 1950

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
1955 1960 1965

Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
1970 1975 1980

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser

1985 1990 1995

Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
2000 2005 2010

Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
2015 2020 2025

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
2030 2035 2040

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser
2045 2050 2055

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
2060 2065 2070

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
2075 2080 2085

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser
2090 2095 2100

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu
2105 2110 2115

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
2120 2125 2130

Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
2135 2140 2145

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
2150 2155 2160

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Ala Ser Ser Glu Gln
2165 2170 2175

Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala Thr Gly His His
2180 2185 2190

His His His His
2195

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<211> 6594
<212> DNA
<213> Artificial Sequence

<220>
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gccacagggt cgccgggatc gtcgactcca tcaggcgcaa cgggatcgcc tggatcaagc 180

actccatccg gagcgacggg aagcccaggg agctccccgt ccgcatcgac tgggacggga 240

cctggagcaa gccccggtac gtcgtcaaca ggttcgcctg gtgcctcgcc tggcacgtcc 300

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ccgggcacct catcaacggg atcaccaggg gcttcaccgg gaacgagctc gacgggttca 420

ccgggagcgt ccccggggac ttcgtcgaca ggtagcccag gggcaagccc cggtacatcg 480

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<211> 2197
<212> PRT
<213> Artificial Sequence

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

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His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
               165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
               245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
               325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
               405                 410                 415

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser
            420                 425                 430
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Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser
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Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr
    450                 455                 460

Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
465                 470                 475                 480

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly
                485                 490                 495

Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
            500                 505                 510

Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
        515                 520                 525

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
    530                 535                 540

Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
545                 550                 555                 560

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr
                565                 570                 575

Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser
            580                 585                 590

Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
        595                 600                 605

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser
    610                 615                 620

Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala
625                 630                 635                 640

Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr
                645                 650                 655

Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr
            660                 665                 670

Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser
        675                 680                 685

Gly Ala Thr Gly Ser Pro Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
    690                 695                 700
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Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
705                 710                 715                 720

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
                725                 730                 735

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
            740                 745                 750

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
        755                 760                 765

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
    770                 775                 780

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
785                 790                 795                 800

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
                805                 810                 815

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
            820                 825                 830

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
        835                 840                 845

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
    850                 855                 860

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
865                 870                 875                 880

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
                885                 890                 895

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
            900                 905                 910

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
        915                 920                 925

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
    930                 935                 940

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
945                 950                 955                 960

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
                965                 970                 975
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Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
            980                 985                 990

Asn Arg Gly Met Thr Ala Leu Leu  Lys Val Ser Ser Cys  Asp Lys Asn
        995                 1000                1005

Thr Gly  Asp Tyr Tyr Glu Asp  Ser Tyr Glu Asp Ile  Ser Ala Tyr
    1010                1015                1020

Leu Leu  Ser Lys Asn Asn Ala  Ile Glu Pro Arg Ser  Phe Ser Gln
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Asn Gly  Ala Pro Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
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Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
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Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1070                1075                1080

Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
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Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
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Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1115                1120                1125

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Thr Glu Pro
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Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1145                1150                1155

Ser Ala  Pro Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1160                1165                1170

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Ala Ser
    1175                1180                1185

Ser Pro  Pro Val Leu Lys Arg  His Gln Ala Glu Ile  Thr Arg Thr
    1190                1195                1200

Thr Leu  Gln Ser Asp Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile
    1205                1210                1215

Ser Val  Glu Met Lys Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp
    1220                1225                1230
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Glu Asn  Gln Ser Pro Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr
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Phe Ile  Ala Ala Val Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser
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Ser Pro  His Val Leu Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro
    1265                 1270                 1275

Gln Phe  Lys Lys Val Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe
    1280                 1285                 1290

Thr Gln  Pro Leu Tyr Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu
    1295                 1300                 1305

Leu Gly  Pro Tyr Ile Arg Ala  Glu Val Glu Asp Asn  Ile Met Val
    1310                 1315                 1320

Thr Phe  Arg Asn Gln Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser
    1325                 1330                 1335

Leu Ile  Ser Tyr Glu Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg
    1340                 1345                 1350

Lys Asn  Phe Val Lys Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys
    1355                 1360                 1365

Val Gln  His His Met Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys
    1370                 1375                 1380

Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu Glu Lys  Asp Val His
    1385                 1390                 1395

Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys His Thr  Asn Thr Leu
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Asn Pro  Ala His Gly Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu
    1415                 1420                 1425

Phe Phe  Thr Ile Phe Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu
    1430                 1435                 1440

Asn Met  Glu Arg Asn Cys Arg  Ala Pro Cys Asn Ile  Gln Met Glu
    1445                 1450                 1455

Asp Pro  Thr Phe Lys Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly
    1460                 1465                 1470

Tyr Ile  Met Asp Thr Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln
    1475                 1480                 1485
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Arg Ile  Arg Trp Tyr Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile
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His Ser  Ile His Phe Ser Gly  His Val Phe Thr Val  Arg Lys Lys
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Glu Glu  Tyr Lys Met Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe
    1520                 1525                 1530

Glu Thr  Val Glu Met Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val
    1535                 1540                 1545

Glu Cys  Leu Ile Gly Glu His  Leu His Ala Gly Met  Ser Thr Leu
    1550                 1555                 1560

Phe Leu  Val Tyr Ser Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala
    1565                 1570                 1575

Ser Gly  His Ile Arg Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr
    1580                 1585                 1590

Gly Gln  Trp Ala Pro Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser
    1595                 1600                 1605

Ile Asn  Ala Trp Ser Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val
    1610                 1615                 1620

Asp Leu  Leu Ala Pro Met Ile  Ile His Gly Ile Lys  Thr Gln Gly
    1625                 1630                 1635

Ala Arg  Gln Lys Phe Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile
    1640                 1645                 1650

Met Tyr  Ser Leu Asp Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn
    1655                 1660                 1665

Ser Thr  Gly Thr Leu Met Val  Phe Phe Gly Asn Val  Asp Ser Ser
    1670                 1675                 1680

Gly Ile  Lys His Asn Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr
    1685                 1690                 1695

Ile Arg  Leu His Pro Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg
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Met Glu  Leu Met Gly Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu
    1715                 1720                 1725

Gly Met  Glu Ser Lys Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser
    1730                 1735                 1740
```

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Ser Tyr  Phe Thr Asn Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala
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Arg Leu  His Leu Gln Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val
    1760                1765                1770

Asn Asn  Pro Lys Glu Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met
    1775                1780                1785

Lys Val  Thr Gly Val Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr
    1790                1795                1800

Ser Met  Tyr Val Lys Glu Phe  Leu Ile Ser Ser Ser  Gln Asp Gly
    1805                1810                1815

His Gln  Trp Thr Leu Phe Phe  Gln Asn Gly Lys Val  Lys Val Phe
    1820                1825                1830

Gln Gly  Asn Gln Asp Ser Phe  Thr Pro Val Val Asn  Ser Leu Asp
    1835                1840                1845

Pro Pro  Leu Leu Thr Arg Tyr  Leu Arg Ile His Pro  Gln Ser Trp
    1850                1855                1860

Val His  Gln Ile Ala Leu Arg  Met Glu Val Leu Gly  Cys Glu Ala
    1865                1870                1875

Gln Asp  Leu Tyr Gly Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1880                1885                1890

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1895                1900                1905

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1910                1915                1920

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    1925                1930                1935

Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu
    1940                1945                1950

Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    1955                1960                1965

Glu Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1970                1975                1980

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    1985                1990                1995
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Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
2000 2005 2010

Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
2015 2020 2025

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
2030 2035 2040

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser
2045 2050 2055

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
2060 2065 2070

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
2075 2080 2085

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser
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Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu
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Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
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Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
2135 2140 2145

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
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Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Ala Ser Ser Glu Gln
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Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala Thr Gly His His
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His His His His
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

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Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445
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Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720
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Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
            740                 745                 750

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
        755                 760                 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
    770                 775                 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785                 790                 795                 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                805                 810                 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            820                 825                 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
        835                 840                 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    850                 855                 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865                 870                 875                 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
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Ser Pro Gly Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser
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Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Ser  Ser Thr Pro
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Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
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Gly Thr  Gly Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
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Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Thr
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Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro Gly Ala  Ser Pro Gly
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Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
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Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
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Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Thr
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Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro
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Ser Ser  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
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Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser
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Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Ser  Ser Thr Pro
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Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
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Gly Thr  Gly Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
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Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Thr
    1235                1240                1245
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Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro
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Ser Gly  Ala Thr Gly Ser Pro  Ala Ser Ser Arg Ala  Gln Ser Gly
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Ser Val  Pro Gln Phe Lys Lys  Val Val Phe Gln Glu  Phe Thr Asp
    1280                1285                1290

Gly Ser  Phe Thr Gln Pro Leu  Tyr Arg Gly Glu Leu  Asn Glu His
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Leu Gly  Leu Leu Gly Pro Tyr  Ile Arg Ala Glu Val  Glu Asp Asn
    1310                1315                1320

Ile Met  Val Thr Phe Arg Asn  Gln Ala Ser Arg Pro  Tyr Ser Phe
    1325                1330                1335

Tyr Ser  Ser Leu Ile Ser Tyr  Glu Glu Asp Gln Arg  Gln Gly Ala
    1340                1345                1350

Glu Pro  Arg Lys Asn Phe Val  Lys Pro Asn Glu Thr  Lys Thr Tyr
    1355                1360                1365

Phe Trp  Lys Val Gln His His  Met Ala Pro Thr Lys  Asp Glu Phe
    1370                1375                1380

Asp Cys  Lys Ala Trp Ala Tyr  Phe Ser Asp Val Asp  Leu Glu Lys
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Asp Val  His Ser Gly Leu Ile  Gly Pro Leu Leu Val  Cys His Thr
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Asn Thr  Leu Asn Pro Ala His  Gly Arg Gln Val Thr  Val Gln Glu
    1415                1420                1425

Phe Ala  Leu Phe Phe Thr Ile  Phe Asp Glu Thr Lys  Ser Trp Tyr
    1430                1435                1440

Phe Thr  Glu Asn Met Glu Arg  Asn Cys Arg Ala Pro  Cys Asn Ile
    1445                1450                1455

Gln Met  Glu Asp Pro Thr Phe  Lys Glu Asn Tyr Arg  Phe His Ala
    1460                1465                1470

Ile Asn  Gly Tyr Ile Met Asp  Thr Leu Pro Gly Leu  Val Met Ala
    1475                1480                1485

Gln Asp  Gln Arg Ile Arg Trp  Tyr Leu Leu Ser Met  Gly Ser Asn
    1490                1495                1500
```

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Glu Asn  Ile His Ser Ile His  Phe Ser Gly His Val  Phe Thr Val
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Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
    1520                 1525                 1530

Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    1535                 1540                 1545

Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    1550                 1555                 1560

Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    1565                 1570                 1575

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
    1580                 1585                 1590

Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
    1595                 1600                 1605

Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
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Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
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Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    1640                 1645                 1650

Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
    1655                 1660                 1665

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
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Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
    1685                 1690                 1695

Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1700                 1705                 1710

Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
    1715                 1720                 1725

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
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Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
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```

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Ser Lys  Ala Arg Leu His Leu  Gln Gly Arg Ser Asn  Ala Trp Arg
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Pro Gln  Val Asn Asn Pro Lys  Glu Trp Leu Gln Val  Asp Phe Gln
    1775                1780                1785

Lys Thr  Met Lys Val Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser
    1790                1795                1800

Leu Leu  Thr Ser Met Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser
    1805                1810                1815

Gln Asp  Gly His Gln Trp Thr  Leu Phe Phe Gln Asn  Gly Lys Val
    1820                1825                1830

Lys Val  Phe Gln Gly Asn Gln  Asp Ser Phe Thr Pro  Val Val Asn
    1835                1840                1845

Ser Leu  Asp Pro Pro Leu Leu  Thr Arg Tyr Leu Arg  Ile His Pro
    1850                1855                1860

Gln Ser  Trp Val His Gln Ile  Ala Leu Arg Met Glu  Val Leu Gly
    1865                1870                1875

Cys Glu  Ala Gln Asp Leu Tyr  Gly Ala Pro Gly Thr  Ser Glu Ser
    1880                1885                1890

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1895                1900                1905

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1910                1915                1920

Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr
    1925                1930                1935

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Thr  Ser Thr Glu
    1940                1945                1950

Pro Ser  Glu Gly Ser Ala Pro  Gly Ser Pro Ala Gly  Ser Pro Thr
    1955                1960                1965

Ser Thr  Glu Glu Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1970                1975                1980

Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr
    1985                1990                1995

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Pro Ala Gly
    2000                2005                2010
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Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
2015 2020 2025

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
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Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
2045 2050 2055

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser
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Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
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Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
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Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr
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Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
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Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
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Gly His His His His His His
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

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Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
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Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

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                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
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Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720
```

```
Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Ser Pro Ala Gly
            740                 745                 750

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
        755                 760                 765

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
    770                 775                 780

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
785                 790                 795                 800

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                805                 810                 815

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            820                 825                 830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
        835                 840                 845

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    850                 855                 860

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
865                 870                 875                 880

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990
```

```
Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
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Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
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Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1145                1150                1155

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser
    1160                1165                1170

Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Ser  Ser Thr Pro
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Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1190                1195                1200

Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
    1205                1210                1215

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1220                1225                1230

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Thr  Pro Gly Ser
    1235                1240                1245
```

```
Gly Thr  Ala Ser Ser Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
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Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
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Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1280                 1285                 1290

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Thr  Pro Gly Ser
    1295                 1300                 1305

Gly Thr  Ala Ser Ser Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
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Thr Gly  Ser Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
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Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Thr
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Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro
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Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1370                 1375                 1380

Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
    1385                 1390                 1395

Pro Gly  Ser Ser Pro Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ala
    1400                 1405                 1410

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Thr  Pro Gly Ser
    1415                 1420                 1425

Gly Thr  Ala Ser Ser Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1430                 1435                 1440

Thr Gly  Ser Pro Ala Ser Ser  Ala Pro Cys Asn Ile  Gln Met Glu
    1445                 1450                 1455

Asp Pro  Thr Phe Lys Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly
    1460                 1465                 1470

Tyr Ile  Met Asp Thr Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln
    1475                 1480                 1485

Arg Ile  Arg Trp Tyr Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile
    1490                 1495                 1500
```

```
His Ser  Ile His Phe Ser Gly  His Val Phe Thr Val  Arg Lys Lys
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Glu Glu  Tyr Lys Met Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe
    1520                1525                1530

Glu Thr  Val Glu Met Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val
    1535                1540                1545

Glu Cys  Leu Ile Gly Glu His  Leu His Ala Gly Met  Ser Thr Leu
    1550                1555                1560

Phe Leu  Val Tyr Ser Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala
    1565                1570                1575

Ser Gly  His Ile Arg Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr
    1580                1585                1590

Gly Gln  Trp Ala Pro Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser
    1595                1600                1605

Ile Asn  Ala Trp Ser Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val
    1610                1615                1620

Asp Leu  Leu Ala Pro Met Ile  Ile His Gly Ile Lys  Thr Gln Gly
    1625                1630                1635

Ala Arg  Gln Lys Phe Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile
    1640                1645                1650

Met Tyr  Ser Leu Asp Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn
    1655                1660                1665

Ser Thr  Gly Thr Leu Met Val  Phe Phe Gly Asn Val  Asp Ser Ser
    1670                1675                1680

Gly Ile  Lys His Asn Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr
    1685                1690                1695

Ile Arg  Leu His Pro Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg
    1700                1705                1710

Met Glu  Leu Met Gly Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu
    1715                1720                1725

Gly Met  Glu Ser Lys Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser
    1730                1735                1740

Ser Tyr  Phe Thr Asn Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala
    1745                1750                1755
```

```
Arg Leu  His Leu Gln Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val
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Asn Asn  Pro Lys Glu Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met
    1775                1780                1785

Lys Val  Thr Gly Val Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr
    1790                1795                1800

Ser Met  Tyr Val Lys Glu Phe  Leu Ile Ser Ser Ser  Gln Asp Gly
    1805                1810                1815

His Gln  Trp Thr Leu Phe Phe  Gln Asn Gly Lys Val  Lys Val Phe
    1820                1825                1830

Gln Gly  Asn Gln Asp Ser Phe  Thr Pro Val Val Asn  Ser Leu Asp
    1835                1840                1845

Pro Pro  Leu Leu Thr Arg Tyr  Leu Arg Ile His Pro  Gln Ser Trp
    1850                1855                1860

Val His  Gln Ile Ala Leu Arg  Met Glu Val Leu Gly  Cys Glu Ala
    1865                1870                1875

Gln Asp  Leu Tyr Gly Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1880                1885                1890

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1895                1900                1905

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1910                1915                1920

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    1925                1930                1935

Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu
    1940                1945                1950

Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    1955                1960                1965

Glu Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1970                1975                1980

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    1985                1990                1995

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Pro Ala Gly  Ser Pro Thr
    2000                2005                2010
```

Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
2015 2020 2025

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
2030 2035 2040

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser
2045 2050 2055

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
2060 2065 2070

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
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Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser
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Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu
2105 2110 2115

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
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Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
2135 2140 2145

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
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Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Ala Ser Ser Glu Gln
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His His His His
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Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
35 40 45

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly
50 55 60

Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
65 70 75 80

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
85 90 95

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
100 105 110

Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
115 120 125

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
130 135 140

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
145 150 155 160

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
165 170 175

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro
180 185 190

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Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
    210                 215                 220

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
225                 230                 235                 240

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
                245                 250                 255

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
            260                 265                 270

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
        275                 280                 285

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
    290                 295                 300

Ala Thr Gly Ser Pro Ala Ser Ser Ser Asp Leu Gly Glu Leu Pro Val
305                 310                 315                 320

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
                325                 330                 335

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
            340                 345                 350

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
        355                 360                 365

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
    370                 375                 380

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
385                 390                 395                 400

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
                405                 410                 415

Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
            420                 425                 430

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
        435                 440                 445

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
    450                 455                 460
```

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Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
465                 470                 475                 480

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
                485                 490                 495

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
            500                 505                 510

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
        515                 520                 525

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
    530                 535                 540

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
545                 550                 555                 560

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
                565                 570                 575

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
            580                 585                 590

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
        595                 600                 605

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
    610                 615                 620

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
625                 630                 635                 640

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
                645                 650                 655

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
            660                 665                 670

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
        675                 680                 685

Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
    690                 695                 700

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
705                 710                 715                 720

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
                725                 730                 735
```

```
Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
            740                 745                 750

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
        755                 760                 765

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
    770                 775                 780

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
785                 790                 795                 800

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
                805                 810                 815

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
            820                 825                 830

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
        835                 840                 845

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
    850                 855                 860

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
865                 870                 875                 880

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
                885                 890                 895

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
            900                 905                 910

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
        915                 920                 925

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
    930                 935                 940

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
945                 950                 955                 960

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
                965                 970                 975

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
            980                 985                 990

Asn Arg Gly Met Thr Ala Leu Leu  Lys Val Ser Ser Cys  Asp Lys Asn
        995                 1000                1005
```

```
Thr Gly  Asp Tyr Tyr Glu Asp  Ser Tyr Glu Asp Ile  Ser Ala Tyr
    1010                1015                1020

Leu Leu  Ser Lys Asn Asn Ala  Ile Glu Pro Arg Ser  Phe Ser Gln
    1025                1030                1035

Asn Pro  Pro Val Leu Lys Arg  His Gln Arg Glu Ile  Thr Arg Thr
    1040                1045                1050

Thr Leu  Gln Ser Asp Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile
    1055                1060                1065

Ser Val  Glu Met Lys Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp
    1070                1075                1080

Glu Asn  Gln Ser Pro Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr
    1085                1090                1095

Phe Ile  Ala Ala Val Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser
    1100                1105                1110

Ser Pro  His Val Leu Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro
    1115                1120                1125

Gln Phe  Lys Lys Val Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe
    1130                1135                1140

Thr Gln  Pro Leu Tyr Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu
    1145                1150                1155

Leu Gly  Pro Tyr Ile Arg Ala  Glu Val Glu Asp Asn  Ile Met Val
    1160                1165                1170

Thr Phe  Arg Asn Gln Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser
    1175                1180                1185

Leu Ile  Ser Tyr Glu Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg
    1190                1195                1200

Lys Asn  Phe Val Lys Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys
    1205                1210                1215

Val Gln  His His Met Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys
    1220                1225                1230

Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu Glu Lys  Asp Val His
    1235                1240                1245

Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys His Thr  Asn Thr Leu
    1250                1255                1260
```

```
Asn Pro  Ala His Gly Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu
    1265                 1270                 1275

Phe Phe  Thr Ile Phe Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu
    1280                 1285                 1290

Asn Met  Glu Arg Asn Cys Arg  Ala Pro Cys Asn Ile  Gln Met Glu
    1295                 1300                 1305

Asp Pro  Thr Phe Lys Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly
    1310                 1315                 1320

Tyr Ile  Met Asp Thr Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln
    1325                 1330                 1335

Arg Ile  Arg Trp Tyr Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile
    1340                 1345                 1350

His Ser  Ile His Phe Ser Gly  His Val Phe Thr Val  Arg Lys Lys
    1355                 1360                 1365

Glu Glu  Tyr Lys Met Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe
    1370                 1375                 1380

Glu Thr  Val Glu Met Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val
    1385                 1390                 1395

Glu Cys  Leu Ile Gly Glu His  Leu His Ala Gly Met  Ser Thr Leu
    1400                 1405                 1410

Phe Leu  Val Tyr Ser Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala
    1415                 1420                 1425

Ser Gly  His Ile Arg Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr
    1430                 1435                 1440

Gly Gln  Trp Ala Pro Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser
    1445                 1450                 1455

Ile Asn  Ala Trp Ser Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val
    1460                 1465                 1470

Asp Leu  Leu Ala Pro Met Ile  Ile His Gly Ile Lys  Thr Gln Gly
    1475                 1480                 1485

Ala Arg  Gln Lys Phe Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile
    1490                 1495                 1500

Met Tyr  Ser Leu Asp Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn
    1505                 1510                 1515
```

```
Ser Thr  Gly Thr Leu Met Val  Phe Phe Gly Asn Val  Asp Ser Ser
    1520                1525                1530

Gly Ile  Lys His Asn Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr
    1535                1540                1545

Ile Arg  Leu His Pro Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg
    1550                1555                1560

Met Glu  Leu Met Gly Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu
    1565                1570                1575

Gly Met  Glu Ser Lys Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser
    1580                1585                1590

Ser Tyr  Phe Thr Asn Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala
    1595                1600                1605

Arg Leu  His Leu Gln Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val
    1610                1615                1620

Asn Asn  Pro Lys Glu Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met
    1625                1630                1635

Lys Val  Thr Gly Val Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr
    1640                1645                1650

Ser Met  Tyr Val Lys Glu Phe  Leu Ile Ser Ser Ser  Gln Asp Gly
    1655                1660                1665

His Gln  Trp Thr Leu Phe Phe  Gln Asn Gly Lys Val  Lys Val Phe
    1670                1675                1680

Gln Gly  Asn Gln Asp Ser Phe  Thr Pro Val Val Asn  Ser Leu Asp
    1685                1690                1695

Pro Pro  Leu Leu Thr Arg Tyr  Leu Arg Ile His Pro  Gln Ser Trp
    1700                1705                1710

Val His  Gln Ile Ala Leu Arg  Met Glu Val Leu Gly  Cys Glu Ala
    1715                1720                1725

Gln Asp  Leu Tyr Gly Ala Gly  Ser Pro Gly Ala Glu  Thr Ala Glu
    1730                1735                1740

Gln Lys  Leu Ile Ser Glu Glu  Asp Leu Ser Pro Ala  Thr Gly His
    1745                1750                1755

His His  His His His
    1760
```

<210> 1394
<211> 5292
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polynucleotide

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gccacagggt cgccgggatc gtcgactcca tcaggcgcaa cgggatcgcc tggatcaagc 180

actccatccg gagcgacggg aagcccaggg agctccccgt ccgcatcgac tgggacggga 240

cctggagcaa gccccggtac gtcgtcaaca ggttcgcctg gtgcctcgcc tggcacgtcc 300

agcacgggga gccccggaac gccgggttcc ggaaccgcat cctcctcacc aggtgcctca 360

ccgggcacct catcaacggg atcaccaggg gcttcaccgg gaacgagctc gacgggttca 420

ccgggagcgt ccccggggac ttcgtcgaca ggtagcccag gggcaagccc cggtacatcg 480

tcaacgggtt ccccagggac accggggagc ggtactgcat cgtcctcacc cggttcctcg 540

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gctccctgca atatccagat ggaagatccc acttttaaag agaattatcg cttccatgca 3960

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acggccgaac aaaaactcat ctcagaagag gatctgtcac ctgcaaccgg tcatcatcac 5280

catcaccatt ga 5292

<210> 1395
<211> 1763
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1395
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1 5 10 15

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285
```

```
Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
                405                 410                 415

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser
            420                 425                 430

Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser
        435                 440                 445

Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr
    450                 455                 460

Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
465                 470                 475                 480

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly
                485                 490                 495

Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
            500                 505                 510

Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
        515                 520                 525

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
    530                 535                 540

Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
545                 550                 555                 560
```

```
Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr
               565                 570                 575

Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser
            580                 585                 590

Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
        595                 600                 605

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser
    610                 615                 620

Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala
625                 630                 635                 640

Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr
               645                 650                 655

Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr
            660                 665                 670

Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser
        675                 680                 685

Gly Ala Thr Gly Ser Pro Ala Ser Ser Asp Arg Ser Tyr Lys Ser Gln
    690                 695                 700

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
705                 710                 715                 720

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
               725                 730                 735

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
            740                 745                 750

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
        755                 760                 765

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
    770                 775                 780

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
785                 790                 795                 800

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
               805                 810                 815

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
            820                 825                 830
```

```
Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
        835                 840                 845

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
    850                 855                 860

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
865                 870                 875                 880

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
                885                 890                 895

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
            900                 905                 910

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
        915                 920                 925

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
    930                 935                 940

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
945                 950                 955                 960

Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
                965                 970                 975

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
            980                 985                 990

Asn Arg Gly Met Thr Ala Leu Leu  Lys Val Ser Ser Cys  Asp Lys Asn
        995                 1000                1005

Thr Gly  Asp Tyr Tyr Glu Asp  Ser Tyr Glu Asp Ile  Ser Ala Tyr
    1010                1015                1020

Leu Leu  Ser Lys Asn Asn Ala  Ile Glu Pro Arg Ser  Phe Ser Gln
    1025                1030                1035

Asn Pro  Pro Val Leu Lys Arg  His Gln Arg Glu Ile  Thr Arg Thr
    1040                1045                1050

Thr Leu  Gln Ser Asp Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile
    1055                1060                1065

Ser Val  Glu Met Lys Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp
    1070                1075                1080

Glu Asn  Gln Ser Pro Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr
    1085                1090                1095
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Phe Ile  Ala Ala Val Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser
    1100                1105                1110

Ser Pro  His Val Leu Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro
    1115                1120                1125

Gln Phe  Lys Lys Val Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe
    1130                1135                1140

Thr Gln  Pro Leu Tyr Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu
    1145                1150                1155

Leu Gly  Pro Tyr Ile Arg Ala  Glu Val Glu Asp Asn  Ile Met Val
    1160                1165                1170

Thr Phe  Arg Asn Gln Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser
    1175                1180                1185

Leu Ile  Ser Tyr Glu Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg
    1190                1195                1200

Lys Asn  Phe Val Lys Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys
    1205                1210                1215

Val Gln  His His Met Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys
    1220                1225                1230

Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu Glu Lys  Asp Val His
    1235                1240                1245

Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys His Thr  Asn Thr Leu
    1250                1255                1260

Asn Pro  Ala His Gly Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu
    1265                1270                1275

Phe Phe  Thr Ile Phe Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu
    1280                1285                1290

Asn Met  Glu Arg Asn Cys Arg  Ala Pro Cys Asn Ile  Gln Met Glu
    1295                1300                1305

Asp Pro  Thr Phe Lys Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly
    1310                1315                1320

Tyr Ile  Met Asp Thr Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln
    1325                1330                1335

Arg Ile  Arg Trp Tyr Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile
    1340                1345                1350
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His Ser  Ile His Phe Ser Gly  His Val Phe Thr Val  Arg Lys Lys
    1355                1360                1365

Glu Glu  Tyr Lys Met Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe
    1370                1375                1380

Glu Thr  Val Glu Met Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val
    1385                1390                1395

Glu Cys  Leu Ile Gly Glu His  Leu His Ala Gly Met  Ser Thr Leu
    1400                1405                1410

Phe Leu  Val Tyr Ser Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala
    1415                1420                1425

Ser Gly  His Ile Arg Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr
    1430                1435                1440

Gly Gln  Trp Ala Pro Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser
    1445                1450                1455

Ile Asn  Ala Trp Ser Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val
    1460                1465                1470

Asp Leu  Leu Ala Pro Met Ile  Ile His Gly Ile Lys  Thr Gln Gly
    1475                1480                1485

Ala Arg  Gln Lys Phe Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile
    1490                1495                1500

Met Tyr  Ser Leu Asp Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn
    1505                1510                1515

Ser Thr  Gly Thr Leu Met Val  Phe Phe Gly Asn Val  Asp Ser Ser
    1520                1525                1530

Gly Ile  Lys His Asn Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr
    1535                1540                1545

Ile Arg  Leu His Pro Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg
    1550                1555                1560

Met Glu  Leu Met Gly Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu
    1565                1570                1575

Gly Met  Glu Ser Lys Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser
    1580                1585                1590

Ser Tyr  Phe Thr Asn Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala
    1595                1600                1605
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Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg Pro Gln Val
1610 1615 1620

Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln Lys Thr Met
1625 1630 1635

Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser Leu Leu Thr
1640 1645 1650

Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly
1655 1660 1665

His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val Lys Val Phe
1670 1675 1680

Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn Ser Leu Asp
1685 1690 1695

Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro Gln Ser Trp
1700 1705 1710

Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly Cys Glu Ala
1715 1720 1725

Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu Thr Ala Glu
1730 1735 1740

Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala Thr Gly His
1745 1750 1755

His His His His His
1760

<210> 1396
<211> 5292
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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ctcggtgagc tgcctgtgga cgcaagattt cctcctagag tgccaaaatc ttttccattc 120

aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg gcgcgccagg tacgcccggg tcgggaacag cgtcctcaag cccgggtagc 1260

tcgacgccgt caggagccac agggtcgccg ggatcgtcga ctccatcagg cgcaacggga 1320

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tcacccggtt cctcgacacc cagcggagcc acgggttcgc ccggtactcc ggggtccgga 1740

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tcccctggat cgtcaactcc ctcgggcgcg actgggagcc caggaagctc accgtcggcc 1920

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ggaatcttgg gacctttact ttatggggaa gttggagaca cactgttgat tatatttaag 2280

aatcaagcaa gcagaccata taacatctac cctcacggaa tcactgatgt ccgtcctttg 2340

tattcaagga gattaccaaa aggtgtaaaa catttgaagg attttccaat tctgccagga 2400

gaaatattca aatataaatg gacagtgact gtagaagatg ggccaactaa atcagatcct 2460

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aaggtttcta gttgtgacaa gaacactggt gattattacg aggacagtta tgaagatatt 3060

tcagcatact tgctgagtaa aaacaatgcc attgaaccaa gaagcttctc tcaaaaccca 3120

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gaaatcgatt atgatgatac catatcagtt gaaatgaaga aggaagattt tgacatttat 3240

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gctgcagtgg agaggctctg ggattatggg atgagtagct ccccacatgt tctaagaaac 3360

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ccatatataa gagcagaagt tgaagataat atcatggtaa ctttcagaaa tcaggcctct 3540

cgtccctatt ccttctattc tagccttatt tcttatgagg aagatcagag gcaaggagca 3600

gaacctagaa aaaactttgt caagcctaat gaaaccaaaa cttacttttg gaaagtgcaa 3660

catcatatgg cacccactaa agatgagttt gactgcaaag cctgggctta tttctctgat 3720

gttgacctgg aaaaagatgt gcactcaggc ctgattggac cccttctggt ctgccacact 3780

aacacactga accctgctca tgggagacaa gtgacagtac aggaatttgc tctgtttttc 3840

accatctttg atgagaccaa aagctggtac ttcactgaaa atatggaaag aaactgcagg 3900

gctccctgca atatccagat ggaagatccc acttttaaag agaattatcg cttccatgca 3960

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ggtgtttttg agacagtgga aatgttacca tccaaagctg gaatttggcg ggtggaatgc 4200

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aaccctccaa ttattgctcg atacatccgt ttgcacccaa ctcattatag cattcgcagc 4680

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cctcaggtga ataatccaaa agagtggctg caagtggact tccagaagac aatgaaagtc 4920

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ttactgactc gctaccttcg aattcacccc cagagttggg tgcaccagat tgccctgagg 5160

atggaggttc tgggctgcga ggcacaggac ctctacggcg ccggatcacc tggggccgaa 5220

acggccgaac aaaaactcat ctcagaagag gatctgtcac ctgcaaccgg tcatcatcac 5280

catcaccatt ga 5292

<210> 1397
<211> 1766
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1397

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val

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        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
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385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro

660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
740 745 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
755 760 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
770 775 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785 790 795 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
805 810 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Gly Ala Pro
820 825 830

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
835 840 845

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
850 855 860

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
865 870 875 880

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
885 890 895

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
900 905 910

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
915 920 925

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro

930 935 940

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
945 950 955 960

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
965 970 975

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
980 985 990

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
995 1000 1005

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
1010 1015 1020

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser
1025 1030 1035

Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
1040 1045 1050

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
1055 1060 1065

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
1070 1075 1080

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr
1085 1090 1095

Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
1100 1105 1110

Ser Gly Ala Thr Gly Ser Pro Ala Ser Ser Arg Ala Gln Ser Gly
1115 1120 1125

Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp
1130 1135 1140

Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His
1145 1150 1155

Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn
1160 1165 1170

Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe
1175 1180 1185

Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala

1190 1195 1200

Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr Lys Thr Tyr
1205 1210 1215

Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp Glu Phe
1220 1225 1230

Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu Lys
1235 1240 1245

Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val Cys His Thr
1250 1255 1260

Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr Val Gln Glu
1265 1270 1275

Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys Ser Trp Tyr
1280 1285 1290

Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro Cys Asn Ile
1295 1300 1305

Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg Phe His Ala
1310 1315 1320

Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu Val Met Ala
1325 1330 1335

Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met Gly Ser Asn
1340 1345 1350

Glu Asn Ile His Ser Ile His Phe Ser Gly His Val Phe Thr Val
1355 1360 1365

Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn Leu Tyr Pro
1370 1375 1380

Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys Ala Gly Ile
1385 1390 1395

Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His Ala Gly Met
1400 1405 1410

Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln Thr Pro Leu
1415 1420 1425

Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile Thr Ala Ser
1430 1435 1440

Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg Leu His Tyr

1445 1450 1455

Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro Phe Ser Trp
1460 1465 1470

Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His Gly Ile Lys
1475 1480 1485

Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr Ile Ser Gln
1490 1495 1500

Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp Gln Thr Tyr
1505 1510 1515

Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe Gly Asn Val
1520 1525 1530

Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro Pro Ile Ile
1535 1540 1545

Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser Ile Arg Ser
1550 1555 1560

Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn Ser Cys Ser
1565 1570 1575

Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp Ala Gln Ile
1580 1585 1590

Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp Ser Pro
1595 1600 1605

Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg
1610 1615 1620

Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln
1625 1630 1635

Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser
1640 1645 1650

Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser
1655 1660 1665

Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val
1670 1675 1680

Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn
1685 1690 1695

Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro

1700 1705 1710

Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly
1715 1720 1725

Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu
1730 1735 1740

Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala
1745 1750 1755

Thr Gly His His His His His His
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atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

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<210> 1399
<211> 1763
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

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Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495
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His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765
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Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
        995                 1000                1005

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser
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Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Ser  Ser Thr Pro
    1025                1030                1035
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Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
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Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
    1055                1060                1065

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1070                1075                1080

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Thr  Pro Gly Ser
    1085                1090                1095

Gly Thr  Ala Ser Ser Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1100                1105                1110

Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    1115                1120                1125

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1130                1135                1140

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Thr  Pro Gly Ser
    1145                1150                1155

Gly Thr  Ala Ser Ser Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1160                1165                1170

Thr Gly  Ser Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1175                1180                1185

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Thr
    1190                1195                1200

Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro
    1205                1210                1215

Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1220                1225                1230

Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
    1235                1240                1245

Pro Gly  Ser Ser Pro Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ala
    1250                1255                1260

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Thr  Pro Gly Ser
    1265                1270                1275

Gly Thr  Ala Ser Ser Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1280                1285                1290
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Thr Gly  Ser Pro Ala Ser Ser  Ala Pro Cys Asn Ile  Gln Met Glu
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Asp Pro  Thr Phe Lys Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly
    1310                1315                1320

Tyr Ile  Met Asp Thr Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln
    1325                1330                1335

Arg Ile  Arg Trp Tyr Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile
    1340                1345                1350

His Ser  Ile His Phe Ser Gly  His Val Phe Thr Val  Arg Lys Lys
    1355                1360                1365

Glu Glu  Tyr Lys Met Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe
    1370                1375                1380

Glu Thr  Val Glu Met Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val
    1385                1390                1395

Glu Cys  Leu Ile Gly Glu His  Leu His Ala Gly Met  Ser Thr Leu
    1400                1405                1410

Phe Leu  Val Tyr Ser Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala
    1415                1420                1425

Ser Gly  His Ile Arg Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr
    1430                1435                1440

Gly Gln  Trp Ala Pro Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser
    1445                1450                1455

Ile Asn  Ala Trp Ser Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val
    1460                1465                1470

Asp Leu  Leu Ala Pro Met Ile  Ile His Gly Ile Lys  Thr Gln Gly
    1475                1480                1485

Ala Arg  Gln Lys Phe Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile
    1490                1495                1500

Met Tyr  Ser Leu Asp Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn
    1505                1510                1515

Ser Thr  Gly Thr Leu Met Val  Phe Phe Gly Asn Val  Asp Ser Ser
    1520                1525                1530

Gly Ile  Lys His Asn Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr
    1535                1540                1545
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Ile Arg Leu His Pro Thr His Tyr Ser Ile Arg Ser Thr Leu Arg
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Met Glu Leu Met Gly Cys Asp Leu Asn Ser Cys Ser Met Pro Leu
1565 1570 1575

Gly Met Glu Ser Lys Ala Ile Ser Asp Ala Gln Ile Thr Ala Ser
1580 1585 1590

Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp Ser Pro Ser Lys Ala
1595 1600 1605

Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg Pro Gln Val
1610 1615 1620

Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln Lys Thr Met
1625 1630 1635

Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser Leu Leu Thr
1640 1645 1650

Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly
1655 1660 1665

His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val Lys Val Phe
1670 1675 1680

Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn Ser Leu Asp
1685 1690 1695

Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro Gln Ser Trp
1700 1705 1710

Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly Cys Glu Ala
1715 1720 1725

Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu Thr Ala Glu
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Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala Thr Gly His
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His His His His His His
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<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic polynucleotide

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acggccgaac aaaaactcat ctcagaagag gatctgtcac ctgcaaccgg tcatcatcac 5280

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<211> 1501
<212> PRT
<213> Artificial Sequence

<220>
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

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Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320
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Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590
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Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860
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Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1025                1030                1035

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1040                1045                1050

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1055                1060                1065

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1070                1075                1080

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1085                1090                1095

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1100                1105                1110

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1115                1120                1125
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Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1130                1135                1140

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1145                1150                1155

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1160                1165                1170

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1175                1180                1185

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1190                1195                1200

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1205                1210                1215

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1220                1225                1230

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1235                1240                1245

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1250                1255                1260

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1265                1270                1275

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1280                1285                1290

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1295                1300                1305

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1310                1315                1320

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1325                1330                1335

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1340                1345                1350

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1355                1360                1365

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1370                1375                1380
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Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1385 1390 1395

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1400 1405 1410

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1415 1420 1425

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Pro Gly Ser
1430 1435 1440

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
1445 1450 1455

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
1460 1465 1470

Ser Glu Thr Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu
1475 1480 1485

Asp Leu Ser Pro Ala Thr Gly His His His His His His
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<210> 1402
<211> 4506
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

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<211> 1553
<212> PRT
<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

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Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540
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Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Ser Pro Ala
            740                 745                 750

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
        755                 760                 765

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
    770                 775                 780

Ala Ser Ser Pro Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg
785                 790                 795                 800

Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
                805                 810                 815
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Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu
            820                 825                 830

Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile
        835                 840                 845

Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His
    850                 855                 860

Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys
865                 870                 875                 880

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
                885                 890                 895

Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
            900                 905                 910

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser
        915                 920                 925

Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln
    930                 935                 940

Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
945                 950                 955                 960

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp
                965                 970                 975

Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu
            980                 985                 990

Lys Asp Val His Ser Gly Leu Ile  Gly Pro Leu Leu Val  Cys His Thr
        995                 1000                1005

Asn Thr  Leu Asn Pro Ala His  Gly Arg Gln Val Thr  Val Gln Glu
    1010                1015                1020

Phe Ala  Leu Phe Phe Thr Ile  Phe Asp Glu Thr Lys  Ser Trp Tyr
    1025                1030                1035

Phe Thr  Glu Asn Met Glu Arg  Asn Cys Arg Gly Ala  Pro Gly Ser
    1040                1045                1050

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    1055                1060                1065

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1070                1075                1080
```

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Ser Glu  Thr Pro Ala Ser Ser  Ala Pro Cys Asn Ile  Gln Met Glu
    1085                 1090                 1095

Asp Pro  Thr Phe Lys Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly
    1100                 1105                 1110

Tyr Ile  Met Asp Thr Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln
    1115                 1120                 1125

Arg Ile  Arg Trp Tyr Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile
    1130                 1135                 1140

His Ser  Ile His Phe Ser Gly  His Val Phe Thr Val  Arg Lys Lys
    1145                 1150                 1155

Glu Glu  Tyr Lys Met Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe
    1160                 1165                 1170

Glu Thr  Val Glu Met Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val
    1175                 1180                 1185

Glu Cys  Leu Ile Gly Glu His  Leu His Ala Gly Met  Ser Thr Leu
    1190                 1195                 1200

Phe Leu  Val Tyr Ser Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala
    1205                 1210                 1215

Ser Gly  His Ile Arg Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr
    1220                 1225                 1230

Gly Gln  Trp Ala Pro Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser
    1235                 1240                 1245

Ile Asn  Ala Trp Ser Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val
    1250                 1255                 1260

Asp Leu  Leu Ala Pro Met Ile  Ile His Gly Ile Lys  Thr Gln Gly
    1265                 1270                 1275

Ala Arg  Gln Lys Phe Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile
    1280                 1285                 1290

Met Tyr  Ser Leu Asp Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn
    1295                 1300                 1305

Ser Thr  Gly Thr Leu Met Val  Phe Phe Gly Asn Val  Asp Ser Ser
    1310                 1315                 1320

Gly Ile  Lys His Asn Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr
    1325                 1330                 1335
```

Ile Arg Leu His Pro Thr His Tyr Ser Ile Arg Ser Thr Leu Arg
1340 1345 1350

Met Glu Leu Met Gly Cys Asp Leu Asn Ser Cys Ser Met Pro Leu
1355 1360 1365

Gly Met Glu Ser Lys Ala Ile Ser Asp Ala Gln Ile Thr Ala Ser
1370 1375 1380

Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp Ser Pro Ser Lys Ala
1385 1390 1395

Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg Pro Gln Val
1400 1405 1410

Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln Lys Thr Met
1415 1420 1425

Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser Leu Leu Thr
1430 1435 1440

Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly
1445 1450 1455

His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val Lys Val Phe
1460 1465 1470

Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn Ser Leu Asp
1475 1480 1485

Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro Gln Ser Trp
1490 1495 1500

Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly Cys Glu Ala
1505 1510 1515

Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu Thr Ala Glu
1520 1525 1530

Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala Thr Gly His
1535 1540 1545

His His His His His His
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<210> 1404
<211> 4662
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

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tccaaagctg gaatttggcg ggtggaatgc cttattggcg agcatctaca tgctgggatg 3600

agcacacttt ttctggtgta cagcaataag tgtcagactc ccctgggaat ggcttctgga 3660

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cagagttggg tgcaccagat tgccctgagg atggaggttc tgggctgcga ggcacaggac 4560

ctctacggcg ccggatcacc tggggccgaa acggccgaac aaaaactcat ctcagaagag 4620

gatctgtcac ctgcaaccgg tcatcatcac catcaccatt ga 4662

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<211> 1661
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1405

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn

130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro

405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp

675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Ser Pro Ala
740 745 750

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
755 760 765

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
770 775 780

Ala Ser Ser Pro Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg
785 790 795 800

Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
805 810 815

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu
820 825 830

Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile
835 840 845

Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His
850 855 860

Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys
865 870 875 880

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
885 890 895

Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
900 905 910

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser
915 920 925

Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln
930 935 940

Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr

945 950 955 960

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp
965 970 975

Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu
980 985 990

Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val Cys His Thr
995 1000 1005

Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr Val Gln Glu
1010 1015 1020

Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys Ser Trp Tyr
1025 1030 1035

Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Gly Ala Pro Gly Thr
1040 1045 1050

Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly
1055 1060 1065

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
1070 1075 1080

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr
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Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly
1115 1120 1125

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
1130 1135 1140

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
1145 1150 1155

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser
1160 1165 1170

Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly
1175 1180 1185

Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Ala Pro Cys Asn Ile
1190 1195 1200

Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg Phe His Ala

1205 1210 1215

Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu Val Met Ala
1220 1225 1230

Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met Gly Ser Asn
1235 1240 1245

Glu Asn Ile His Ser Ile His Phe Ser Gly His Val Phe Thr Val
1250 1255 1260

Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn Leu Tyr Pro
1265 1270 1275

Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys Ala Gly Ile
1280 1285 1290

Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His Ala Gly Met
1295 1300 1305

Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln Thr Pro Leu
1310 1315 1320

Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile Thr Ala Ser
1325 1330 1335

Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg Leu His Tyr
1340 1345 1350

Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro Phe Ser Trp
1355 1360 1365

Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His Gly Ile Lys
1370 1375 1380

Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr Ile Ser Gln
1385 1390 1395

Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp Gln Thr Tyr
1400 1405 1410

Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe Gly Asn Val
1415 1420 1425

Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro Pro Ile Ile
1430 1435 1440

Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser Ile Arg Ser
1445 1450 1455

Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn Ser Cys Ser

1460 1465 1470

Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp Ala Gln Ile
1475 1480 1485

Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp Ser Pro
1490 1495 1500

Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg
1505 1510 1515

Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln
1520 1525 1530

Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser
1535 1540 1545

Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser
1550 1555 1560

Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val
1565 1570 1575

Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn
1580 1585 1590

Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro
1595 1600 1605

Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly
1610 1615 1620

Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu
1625 1630 1635

Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala
1640 1645 1650

Thr Gly His His His His His His
1655 1660

<210> 1406
<211> 4986
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

agagatctag cttcaggact cattggccct ctcctcatct gctacaaaga atctgtagat 1680

caaagaggaa accagataat gtcagacaag aggaatgtca tcctgttttc tgtatttgat 1740

gagaaccgaa gctggtacct cacagagaat atacaacgct ttctccccaa tccagctgga 1800

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gtttttgata gtttgcagtt gtcagtttgt ttgcatgagg tggcatactg gtacattcta 1920

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gacagttatg aagatatttc agcatacttg ctgagtaaaa acaatgccat tgaaccaaga 2220

agcttctctc aaaacggcgc gccaggttct cctgctggct cccccacctc aacagaagag 2280

gggacaagcg aaagcgctac gcctgagagt ggccctggct ctgagccagc cacctccggc 2340

tctgaaaccc ctgcctcgag cccaccagtc ttgaaacgcc atcaagctga aataactcgt 2400

actactcttc agtcagatca agaggaaatc gattatgatg ataccatatc agttgaaatg 2460

aagaaggaag attttgacat ttatgatgag gatgaaaatc agagcccccg cagctttcaa 2520

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gttgttttcc aggaatttac tgatggctcc tttactcagc ccttataccg tggagaacta 2700

aatgaacatt tgggactcct ggggccatat ataagagcag aagttgaaga taatatcatg 2760

gtaactttca gaaatcaggc ctctcgtccc tattccttct attctagcct tatttcttat 2820

gaggaagatc agaggcaagg agcagaacct agaaaaaact ttgtcaagcc taatgaaacc 2880

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acgtcctcaa caggctcgcc cggtgcgtca cccggcacgt cgtccacggg ttcacctggt 3300

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atcatgtata gtcttgatgg gaagaagtgg cagacttatc gaggaaattc cactggaacc 4260

ttaatggtct tctttggcaa tgtggattca tctgggataa aacacaatat ttttaaccct 4320

ccaattattg ctcgatacat ccgtttgcac ccaactcatt atagcattcg cagcactctt 4380

cgcatggagt tgatgggctg tgatttaaat agttgcagca tgccattggg aatggagagt 4440

aaagcaatat cagatgcaca gattactgct tcatcctact ttaccaatat gtttgccacc 4500

tggtctcctt caaaagctcg acttcacctc caagggagga gtaatgcctg gagacctcag 4560

gtgaataatc caaaagagtg gctgcaagtg gacttccaga agacaatgaa agtcacagga 4620

gtaactactc agggagtaaa atctctgctt accagcatgt atgtgaagga gttcctcatc 4680

tccagcagtc aagatggcca tcagtggact ctctttttc agaatggcaa agtaaaggtt 4740

tttcagggaa atcaagactc cttcacacct gtggtgaact ctctagaccc accgttactg 4800

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gttctgggct gcgaggcaca ggacctctac ggcgccggat cacctggggc cgaaacggcc 4920

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cattga 4986

<210> 1407
<211> 1805
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1407
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

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Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380
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Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655
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Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Ser Pro Ala
            740                 745                 750

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
        755                 760                 765

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
    770                 775                 780

Ala Ser Ser Pro Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg
785                 790                 795                 800

Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
                805                 810                 815

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu
            820                 825                 830

Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile
        835                 840                 845

Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His
    850                 855                 860

Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys
865                 870                 875                 880

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
                885                 890                 895

Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
            900                 905                 910

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser
        915                 920                 925
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Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln
    930                 935                 940

Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
945                 950                 955                 960

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp
                965                 970                 975

Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu
            980                 985                 990

Lys Asp Val His Ser Gly Leu Ile  Gly Pro Leu Leu Val  Cys His Thr
        995                 1000                1005

Asn Thr  Leu Asn Pro Ala His  Gly Arg Gln Val Thr  Val Gln Glu
    1010                1015                1020

Phe Ala  Leu Phe Phe Thr Ile  Phe Asp Glu Thr Lys  Ser Trp Tyr
    1025                1030                1035

Phe Thr  Glu Asn Met Glu Arg  Asn Cys Arg Gly Ala  Pro Gly Thr
    1040                1045                1050

Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro
    1055                1060                1065

Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1070                1075                1080

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1085                1090                1095

Pro Gly  Ser Ser Pro Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ala
    1100                1105                1110

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1115                1120                1125

Thr Ser  Ser Thr Gly Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala
    1130                1135                1140

Ser Ser  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    1145                1150                1155

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1160                1165                1170

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1175                1180                1185
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Thr Ser  Ser Thr Gly Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala
    1190                 1195                 1200

Ser Ser  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1205                 1210                 1215

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser
    1220                 1225                 1230

Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Thr  Pro Gly Ser
    1235                 1240                 1245

Gly Thr  Ala Ser Ser Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1250                 1255                 1260

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1265                 1270                 1275

Pro Gly  Ser Ser Pro Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ser
    1280                 1285                 1290

Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro Gly Ala  Ser Pro Gly
    1295                 1300                 1305

Thr Ser  Ser Thr Gly Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala
    1310                 1315                 1320

Ser Ser  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1325                 1330                 1335

Pro Ala  Ser Ser Ala Pro Cys  Asn Ile Gln Met Glu  Asp Pro Thr
    1340                 1345                 1350

Phe Lys  Glu Asn Tyr Arg Phe  His Ala Ile Asn Gly  Tyr Ile Met
    1355                 1360                 1365

Asp Thr  Leu Pro Gly Leu Val  Met Ala Gln Asp Gln  Arg Ile Arg
    1370                 1375                 1380

Trp Tyr  Leu Leu Ser Met Gly  Ser Asn Glu Asn Ile  His Ser Ile
    1385                 1390                 1395

His Phe  Ser Gly His Val Phe  Thr Val Arg Lys Lys  Glu Glu Tyr
    1400                 1405                 1410

Lys Met  Ala Leu Tyr Asn Leu  Tyr Pro Gly Val Phe  Glu Thr Val
    1415                 1420                 1425

Glu Met  Leu Pro Ser Lys Ala  Gly Ile Trp Arg Val  Glu Cys Leu
    1430                 1435                 1440
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Ile Gly  Glu His Leu His Ala  Gly Met Ser Thr Leu  Phe Leu Val
    1445                1450                1455

Tyr Ser  Asn Lys Cys Gln Thr  Pro Leu Gly Met Ala  Ser Gly His
    1460                1465                1470

Ile Arg  Asp Phe Gln Ile Thr  Ala Ser Gly Gln Tyr  Gly Gln Trp
    1475                1480                1485

Ala Pro  Lys Leu Ala Arg Leu  His Tyr Ser Gly Ser  Ile Asn Ala
    1490                1495                1500

Trp Ser  Thr Lys Glu Pro Phe  Ser Trp Ile Lys Val  Asp Leu Leu
    1505                1510                1515

Ala Pro  Met Ile Ile His Gly  Ile Lys Thr Gln Gly  Ala Arg Gln
    1520                1525                1530

Lys Phe  Ser Ser Leu Tyr Ile  Ser Gln Phe Ile Ile  Met Tyr Ser
    1535                1540                1545

Leu Asp  Gly Lys Lys Trp Gln  Thr Tyr Arg Gly Asn  Ser Thr Gly
    1550                1555                1560

Thr Leu  Met Val Phe Phe Gly  Asn Val Asp Ser Ser  Gly Ile Lys
    1565                1570                1575

His Asn  Ile Phe Asn Pro Pro  Ile Ile Ala Arg Tyr  Ile Arg Leu
    1580                1585                1590

His Pro  Thr His Tyr Ser Ile  Arg Ser Thr Leu Arg  Met Glu Leu
    1595                1600                1605

Met Gly  Cys Asp Leu Asn Ser  Cys Ser Met Pro Leu  Gly Met Glu
    1610                1615                1620

Ser Lys  Ala Ile Ser Asp Ala  Gln Ile Thr Ala Ser  Ser Tyr Phe
    1625                1630                1635

Thr Asn  Met Phe Ala Thr Trp  Ser Pro Ser Lys Ala  Arg Leu His
    1640                1645                1650

Leu Gln  Gly Arg Ser Asn Ala  Trp Arg Pro Gln Val  Asn Asn Pro
    1655                1660                1665

Lys Glu  Trp Leu Gln Val Asp  Phe Gln Lys Thr Met  Lys Val Thr
    1670                1675                1680

Gly Val  Thr Thr Gln Gly Val  Lys Ser Leu Leu Thr  Ser Met Tyr
    1685                1690                1695
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Val Lys Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly His Gln Trp
1700 1705 1710

Thr Leu Phe Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn
1715 1720 1725

Gln Asp Ser Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu
1730 1735 1740

Leu Thr Arg Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln
1745 1750 1755

Ile Ala Leu Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu
1760 1765 1770

Tyr Gly Ala Gly Ser Pro Gly Ala Glu Thr Ala Glu Gln Lys Leu
1775 1780 1785

Ile Ser Glu Glu Asp Leu Ser Pro Ala Thr Gly His His His His
1790 1795 1800

His His
1805

<210> 1408
<211> 5418
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser

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His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
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Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
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His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
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Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
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420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
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Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
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Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala

690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Thr Pro Gly
740 745 750

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
755 760 765

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
770 775 780

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
785 790 795 800

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Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
850 855 860

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865 870 875 880

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Pro
885 890 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
900 905 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
915 920 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
930 935 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945 950 955 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn

965 970 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
980 985 990

Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
995 1000 1005

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1010 1015 1020

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1025 1030 1035

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
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Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
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Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
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Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
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Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
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Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1115 1120 1125

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
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Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Gly Ala
1145 1150 1155

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr
1160 1165 1170

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1175 1180 1185

Thr Ser Gly Ser Glu Thr Pro Ala Ser Ser Ala Pro Cys Asn Ile
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Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg Phe His Ala
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Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu Val Met Ala

1220 1225 1230

Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met Gly Ser Asn
1235 1240 1245

Glu Asn Ile His Ser Ile His Phe Ser Gly His Val Phe Thr Val
1250 1255 1260

Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn Leu Tyr Pro
1265 1270 1275

Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys Ala Gly Ile
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Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His Ala Gly Met
1295 1300 1305

Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln Thr Pro Leu
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Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile Thr Ala Ser
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1340 1345 1350

Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro Phe Ser Trp
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Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His Gly Ile Lys
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Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr Ile Ser Gln
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Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp Gln Thr Tyr
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Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe Gly Asn Val
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Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro Pro Ile Ile
1430 1435 1440

Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser Ile Arg Ser
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Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn Ser Cys Ser
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Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp Ala Gln Ile

1475 1480 1485

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Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg
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Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln
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Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser
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Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro
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Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly
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Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu
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Thr Gly His His His His His His
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ggcatcaaga cccagggtgc ccgtcagaag ttctccagcc tctacatctc tcagtttatc 4200

atcatgtata gtcttgatgg gaagaagtgg cagacttatc gaggaaattc cactggaacc 4260

ttaatggtct tctttggcaa tgtggattca tctgggataa aacacaatat ttttaaccct 4320

ccaattattg ctcgatacat ccgtttgcac ccaactcatt atagcattcg cagcactctt 4380

cgcatggagt tgatgggctg tgatttaaat agttgcagca tgccattggg aatggagagt 4440

aaagcaatat cagatgcaca gattactgct tcatcctact ttaccaatat gtttgccacc 4500

tggtctcctt caaaagctcg acttcacctc caagggagga gtaatgcctg gagacctcag 4560

gtgaataatc caaaagagtg gctgcaagtg gacttccaga agacaatgaa agtcacagga 4620

gtaactactc agggagtaaa atctctgctt accagcatgt atgtgaagga gttcctcatc 4680

tccagcagtc aagatggcca tcagtggact ctctttttc agaatggcaa agtaaaggtt 4740

tttcagggaa atcaagactc cttcacacct gtggtgaact ctctagaccc accgttactg 4800

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gttctgggct gcgaggcaca ggacctctac ggcgccggat cacctggggc cgaaacggcc 4920

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cattga 4986

<210> 1411
<211> 1769
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1411
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

```
Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400
```

```
Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670
```

```
Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Thr Pro Gly
            740                 745                 750

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
        755                 760                 765

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    770                 775                 780

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
785                 790                 795                 800

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
                805                 810                 815

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            820                 825                 830

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
        835                 840                 845

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
    850                 855                 860

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
865                 870                 875                 880

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940
```

```
Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1145                1150                1155

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala
    1160                1165                1170

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1175                1180                1185

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1190                1195                1200
```

```
Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
    1205                 1210                 1215

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala
    1220                 1225                 1230

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1235                 1240                 1245

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1250                 1255                 1260

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1265                 1270                 1275

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ala
    1280                 1285                 1290

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Ala Ser  Ser Ala Pro
    1295                 1300                 1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                 1315                 1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                 1330                 1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                 1345                 1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                 1360                 1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                 1375                 1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                 1390                 1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                 1405                 1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                 1420                 1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                 1435                 1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                 1450                 1455
```

```
Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710
```

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Gly Ser Pro
1730 1735 1740

Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1745 1750 1755

Ser Pro Ala Thr Gly His His His His His His
1760 1765

<210> 1412
<211> 5310
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

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cagcccttat accgtggaga actaaatgaa catttgggac tcctggggcc atatataaga 3060

gcagaagttg aagataatat catggtaact ttcagaaatc aggcctctcg tccctattcc 3120

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<210> 1413
<211> 1913
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1413

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Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
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225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys

```
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Thr Pro Gly
            740                 745                 750

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
        755                 760                 765

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
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770 775 780

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
785 790 795 800

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
805 810 815

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
820 825 830

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
835 840 845

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
850 855 860

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
865 870 875 880

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Pro
885 890 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
900 905 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
915 920 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
930 935 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945 950 955 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
965 970 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
980 985 990

Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
995 1000 1005

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1010 1015 1020

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1025 1030 1035

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg

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Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1145                1150                1155

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser
    1160                1165                1170

Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Ser  Ser Thr Pro
    1175                1180                1185

Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1190                1195                1200

Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
    1205                1210                1215

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1220                1225                1230

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Thr  Pro Gly Ser
    1235                1240                1245

Gly Thr  Ala Ser Ser Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1250                1255                1260

Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    1265                1270                1275

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1280                1285                1290

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Thr  Pro Gly Ser
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Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
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Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
1325 1330 1335

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr
1340 1345 1350

Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
1355 1360 1365

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
1370 1375 1380

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
1385 1390 1395

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala
1400 1405 1410

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser
1415 1420 1425

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
1430 1435 1440

Thr Gly Ser Pro Ala Ser Ser Ala Pro Cys Asn Ile Gln Met Glu
1445 1450 1455

Asp Pro Thr Phe Lys Glu Asn Tyr Arg Phe His Ala Ile Asn Gly
1460 1465 1470

Tyr Ile Met Asp Thr Leu Pro Gly Leu Val Met Ala Gln Asp Gln
1475 1480 1485

Arg Ile Arg Trp Tyr Leu Leu Ser Met Gly Ser Asn Glu Asn Ile
1490 1495 1500

His Ser Ile His Phe Ser Gly His Val Phe Thr Val Arg Lys Lys
1505 1510 1515

Glu Glu Tyr Lys Met Ala Leu Tyr Asn Leu Tyr Pro Gly Val Phe
1520 1525 1530

Glu Thr Val Glu Met Leu Pro Ser Lys Ala Gly Ile Trp Arg Val
1535 1540 1545

Glu Cys Leu Ile Gly Glu His Leu His Ala Gly Met Ser Thr Leu

1550 1555 1560

Phe Leu Val Tyr Ser Asn Lys Cys Gln Thr Pro Leu Gly Met Ala
1565 1570 1575

Ser Gly His Ile Arg Asp Phe Gln Ile Thr Ala Ser Gly Gln Tyr
1580 1585 1590

Gly Gln Trp Ala Pro Lys Leu Ala Arg Leu His Tyr Ser Gly Ser
1595 1600 1605

Ile Asn Ala Trp Ser Thr Lys Glu Pro Phe Ser Trp Ile Lys Val
1610 1615 1620

Asp Leu Leu Ala Pro Met Ile Ile His Gly Ile Lys Thr Gln Gly
1625 1630 1635

Ala Arg Gln Lys Phe Ser Ser Leu Tyr Ile Ser Gln Phe Ile Ile
1640 1645 1650

Met Tyr Ser Leu Asp Gly Lys Lys Trp Gln Thr Tyr Arg Gly Asn
1655 1660 1665

Ser Thr Gly Thr Leu Met Val Phe Phe Gly Asn Val Asp Ser Ser
1670 1675 1680

Gly Ile Lys His Asn Ile Phe Asn Pro Pro Ile Ile Ala Arg Tyr
1685 1690 1695

Ile Arg Leu His Pro Thr His Tyr Ser Ile Arg Ser Thr Leu Arg
1700 1705 1710

Met Glu Leu Met Gly Cys Asp Leu Asn Ser Cys Ser Met Pro Leu
1715 1720 1725

Gly Met Glu Ser Lys Ala Ile Ser Asp Ala Gln Ile Thr Ala Ser
1730 1735 1740

Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp Ser Pro Ser Lys Ala
1745 1750 1755

Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg Pro Gln Val
1760 1765 1770

Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln Lys Thr Met
1775 1780 1785

Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser Leu Leu Thr
1790 1795 1800

Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly

1805 1810 1815

His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val Lys Val Phe
1820 1825 1830

Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn Ser Leu Asp
1835 1840 1845

Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro Gln Ser Trp
1850 1855 1860

Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly Cys Glu Ala
1865 1870 1875

Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu Thr Ala Glu
1880 1885 1890

Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala Thr Gly His
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His His His His His
1910

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

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ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

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gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

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gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

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<211> 1805
<212> PRT
<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

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Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380
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Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655
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Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Pro Gly Ala Ser
            740                 745                 750

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
        755                 760                 765

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
    770                 775                 780

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro
785                 790                 795                 800

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
                805                 810                 815

Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
            820                 825                 830

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
        835                 840                 845

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
    850                 855                 860

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
865                 870                 875                 880

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
                885                 890                 895

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
            900                 905                 910

Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
        915                 920                 925
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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
    930                 935                 940

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
945                 950                 955                 960

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
                965                 970                 975

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
            980                 985                 990

Pro Gly Thr Ser Ser Thr Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser
        995                 1000                1005

Thr Gly  Thr Gly Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
    1010                1015                1020

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Ala Ser
    1025                1030                1035

Ser Pro  Pro Val Leu Lys Arg  His Gln Ala Glu Ile  Thr Arg Thr
    1040                1045                1050

Thr Leu  Gln Ser Asp Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile
    1055                1060                1065

Ser Val  Glu Met Lys Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp
    1070                1075                1080

Glu Asn  Gln Ser Pro Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr
    1085                1090                1095

Phe Ile  Ala Ala Val Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser
    1100                1105                1110

Ser Pro  His Val Leu Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro
    1115                1120                1125

Gln Phe  Lys Lys Val Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe
    1130                1135                1140

Thr Gln  Pro Leu Tyr Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu
    1145                1150                1155

Leu Gly  Pro Tyr Ile Arg Ala  Glu Val Glu Asp Asn  Ile Met Val
    1160                1165                1170

Thr Phe  Arg Asn Gln Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser
    1175                1180                1185
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```
Leu Ile  Ser Tyr Glu Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg
    1190                 1195                 1200

Lys Asn  Phe Val Lys Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys
    1205                 1210                 1215

Val Gln  His His Met Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys
    1220                 1225                 1230

Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu Glu Lys  Asp Val His
    1235                 1240                 1245

Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys His Thr  Asn Thr Leu
    1250                 1255                 1260

Asn Pro  Ala His Gly Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu
    1265                 1270                 1275

Phe Phe  Thr Ile Phe Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu
    1280                 1285                 1290

Asn Met  Glu Arg Asn Cys Arg  Gly Ala Pro Gly Ser  Pro Ala Gly
    1295                 1300                 1305

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1310                 1315                 1320

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1325                 1330                 1335

Pro Ala  Ser Ser Ala Pro Cys  Asn Ile Gln Met Glu  Asp Pro Thr
    1340                 1345                 1350

Phe Lys  Glu Asn Tyr Arg Phe  His Ala Ile Asn Gly  Tyr Ile Met
    1355                 1360                 1365

Asp Thr  Leu Pro Gly Leu Val  Met Ala Gln Asp Gln  Arg Ile Arg
    1370                 1375                 1380

Trp Tyr  Leu Leu Ser Met Gly  Ser Asn Glu Asn Ile  His Ser Ile
    1385                 1390                 1395

His Phe  Ser Gly His Val Phe  Thr Val Arg Lys Lys  Glu Glu Tyr
    1400                 1405                 1410

Lys Met  Ala Leu Tyr Asn Leu  Tyr Pro Gly Val Phe  Glu Thr Val
    1415                 1420                 1425

Glu Met  Leu Pro Ser Lys Ala  Gly Ile Trp Arg Val  Glu Cys Leu
    1430                 1435                 1440
```

```
Ile Gly  Glu His Leu His Ala  Gly Met Ser Thr Leu  Phe Leu Val
    1445                1450                1455

Tyr Ser  Asn Lys Cys Gln Thr  Pro Leu Gly Met Ala  Ser Gly His
    1460                1465                1470

Ile Arg  Asp Phe Gln Ile Thr  Ala Ser Gly Gln Tyr  Gly Gln Trp
    1475                1480                1485

Ala Pro  Lys Leu Ala Arg Leu  His Tyr Ser Gly Ser  Ile Asn Ala
    1490                1495                1500

Trp Ser  Thr Lys Glu Pro Phe  Ser Trp Ile Lys Val  Asp Leu Leu
    1505                1510                1515

Ala Pro  Met Ile Ile His Gly  Ile Lys Thr Gln Gly  Ala Arg Gln
    1520                1525                1530

Lys Phe  Ser Ser Leu Tyr Ile  Ser Gln Phe Ile Ile  Met Tyr Ser
    1535                1540                1545

Leu Asp  Gly Lys Lys Trp Gln  Thr Tyr Arg Gly Asn  Ser Thr Gly
    1550                1555                1560

Thr Leu  Met Val Phe Phe Gly  Asn Val Asp Ser Ser  Gly Ile Lys
    1565                1570                1575

His Asn  Ile Phe Asn Pro Pro  Ile Ile Ala Arg Tyr  Ile Arg Leu
    1580                1585                1590

His Pro  Thr His Tyr Ser Ile  Arg Ser Thr Leu Arg  Met Glu Leu
    1595                1600                1605

Met Gly  Cys Asp Leu Asn Ser  Cys Ser Met Pro Leu  Gly Met Glu
    1610                1615                1620

Ser Lys  Ala Ile Ser Asp Ala  Gln Ile Thr Ala Ser  Ser Tyr Phe
    1625                1630                1635

Thr Asn  Met Phe Ala Thr Trp  Ser Pro Ser Lys Ala  Arg Leu His
    1640                1645                1650

Leu Gln  Gly Arg Ser Asn Ala  Trp Arg Pro Gln Val  Asn Asn Pro
    1655                1660                1665

Lys Glu  Trp Leu Gln Val Asp  Phe Gln Lys Thr Met  Lys Val Thr
    1670                1675                1680

Gly Val  Thr Thr Gln Gly Val  Lys Ser Leu Leu Thr  Ser Met Tyr
    1685                1690                1695
```

Val Lys Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly His Gln Trp
1700 1705 1710

Thr Leu Phe Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn
1715 1720 1725

Gln Asp Ser Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu
1730 1735 1740

Leu Thr Arg Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln
1745 1750 1755

Ile Ala Leu Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu
1760 1765 1770

Tyr Gly Ala Gly Ser Pro Gly Ala Glu Thr Ala Glu Gln Lys Leu
1775 1780 1785

Ile Ser Glu Glu Asp Leu Ser Pro Ala Thr Gly His His His His
1790 1795 1800

His His
1805

<210> 1416
<211> 5418
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

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atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg      960
gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat     1020
gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt     1080
gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa     1140
acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc     1200
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gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca     1380
ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc     1440
actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat     1500
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aaaatggtct atgaagacac actcacccta ttcccattct caggagaaac tgtcttcatg     2040
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gcgtcgtcga gcccagggtc gtccacccca agcggggcta ctgggtcccc gggtagctcg     2760
ccgagcgcat caacgggaac cggaccgggc agctcccccct cagcgtcgac agggacaggc     2820
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gaaatcgatt atgatgatac catatcagtt gaaatgaaga aggaagattt tgacatttat 3240

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<210> 1417
<211> 1913
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1417

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser

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145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
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420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala

690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Pro Gly Ala Ser
740 745 750

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
755 760 765

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
770 775 780

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro
785 790 795 800

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
805 810 815

Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
820 825 830

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
835 840 845

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
850 855 860

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
865 870 875 880

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
885 890 895

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
900 905 910

Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
915 920 925

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
930 935 940

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
945 950 955 960

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser

965 970 975

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
980 985 990

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
995 1000 1005

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
1010 1015 1020

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Ala Ser
1025 1030 1035

Ser Pro Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr
1040 1045 1050

Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
1055 1060 1065

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp
1070 1075 1080

Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr
1085 1090 1095

Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser
1100 1105 1110

Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro
1115 1120 1125

Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe
1130 1135 1140

Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
1145 1150 1155

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val
1160 1165 1170

Thr Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser
1175 1180 1185

Leu Ile Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg
1190 1195 1200

Lys Asn Phe Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys
1205 1210 1215

Val Gln His His Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys

1220 1225 1230

Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu Lys Asp Val His
1235 1240 1245

Ser Gly Leu Ile Gly Pro Leu Leu Val Cys His Thr Asn Thr Leu
1250 1255 1260

Asn Pro Ala His Gly Arg Gln Val Thr Val Gln Glu Phe Ala Leu
1265 1270 1275

Phe Phe Thr Ile Phe Asp Glu Thr Lys Ser Trp Tyr Phe Thr Glu
1280 1285 1290

Asn Met Glu Arg Asn Cys Arg Gly Ala Pro Gly Thr Pro Gly Ser
1295 1300 1305

Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
1310 1315 1320

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
1325 1330 1335

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser
1340 1345 1350

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser
1355 1360 1365

Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
1370 1375 1380

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
1385 1390 1395

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser
1400 1405 1410

Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
1415 1420 1425

Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
1430 1435 1440

Thr Gly Ser Pro Ala Ser Ser Ala Pro Cys Asn Ile Gln Met Glu
1445 1450 1455

Asp Pro Thr Phe Lys Glu Asn Tyr Arg Phe His Ala Ile Asn Gly
1460 1465 1470

Tyr Ile Met Asp Thr Leu Pro Gly Leu Val Met Ala Gln Asp Gln

1475 1480 1485

Arg Ile Arg Trp Tyr Leu Leu Ser Met Gly Ser Asn Glu Asn Ile
1490 1495 1500

His Ser Ile His Phe Ser Gly His Val Phe Thr Val Arg Lys Lys
1505 1510 1515

Glu Glu Tyr Lys Met Ala Leu Tyr Asn Leu Tyr Pro Gly Val Phe
1520 1525 1530

Glu Thr Val Glu Met Leu Pro Ser Lys Ala Gly Ile Trp Arg Val
1535 1540 1545

Glu Cys Leu Ile Gly Glu His Leu His Ala Gly Met Ser Thr Leu
1550 1555 1560

Phe Leu Val Tyr Ser Asn Lys Cys Gln Thr Pro Leu Gly Met Ala
1565 1570 1575

Ser Gly His Ile Arg Asp Phe Gln Ile Thr Ala Ser Gly Gln Tyr
1580 1585 1590

Gly Gln Trp Ala Pro Lys Leu Ala Arg Leu His Tyr Ser Gly Ser
1595 1600 1605

Ile Asn Ala Trp Ser Thr Lys Glu Pro Phe Ser Trp Ile Lys Val
1610 1615 1620

Asp Leu Leu Ala Pro Met Ile Ile His Gly Ile Lys Thr Gln Gly
1625 1630 1635

Ala Arg Gln Lys Phe Ser Ser Leu Tyr Ile Ser Gln Phe Ile Ile
1640 1645 1650

Met Tyr Ser Leu Asp Gly Lys Lys Trp Gln Thr Tyr Arg Gly Asn
1655 1660 1665

Ser Thr Gly Thr Leu Met Val Phe Phe Gly Asn Val Asp Ser Ser
1670 1675 1680

Gly Ile Lys His Asn Ile Phe Asn Pro Pro Ile Ile Ala Arg Tyr
1685 1690 1695

Ile Arg Leu His Pro Thr His Tyr Ser Ile Arg Ser Thr Leu Arg
1700 1705 1710

Met Glu Leu Met Gly Cys Asp Leu Asn Ser Cys Ser Met Pro Leu
1715 1720 1725

Gly Met Glu Ser Lys Ala Ile Ser Asp Ala Gln Ile Thr Ala Ser

1730 1735 1740

Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp Ser Pro Ser Lys Ala
1745 1750 1755

Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg Pro Gln Val
1760 1765 1770

Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln Lys Thr Met
1775 1780 1785

Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser Leu Leu Thr
1790 1795 1800

Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly
1805 1810 1815

His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val Lys Val Phe
1820 1825 1830

Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn Ser Leu Asp
1835 1840 1845

Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro Gln Ser Trp
1850 1855 1860

Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly Cys Glu Ala
1865 1870 1875

Gln Asp Leu Tyr Gly Ala Gly Ser Pro Gly Ala Glu Thr Ala Glu
1880 1885 1890

Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala Thr Gly His
1895 1900 1905

His His His His His
1910

<210> 1418
<211> 5742
<212> DNA
<213> Artificial Sequence

<220>
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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

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<210> 1419
<211> 2057
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300
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Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575
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Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Pro Gly Ala Ser
            740                 745                 750

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
        755                 760                 765

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
    770                 775                 780

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro
785                 790                 795                 800

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
                805                 810                 815

Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
            820                 825                 830

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
        835                 840                 845
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Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
    850                 855                 860

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
865                 870                 875                 880

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
                885                 890                 895

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
            900                 905                 910

Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
        915                 920                 925

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
    930                 935                 940

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
945                 950                 955                 960

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
                965                 970                 975

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
            980                 985                 990

Pro Gly Thr Ser Ser Thr Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser
        995                 1000                1005

Thr Gly  Thr Gly Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
    1010                1015                1020

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Ala Ser
    1025                1030                1035

Ser Pro  Pro Val Leu Lys Arg  His Gln Ala Glu Ile  Thr Arg Thr
    1040                1045                1050

Thr Leu  Gln Ser Asp Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile
    1055                1060                1065

Ser Val  Glu Met Lys Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp
    1070                1075                1080

Glu Asn  Gln Ser Pro Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr
    1085                1090                1095

Phe Ile  Ala Ala Val Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser
    1100                1105                1110
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Ser Pro  His Val Leu Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro
    1115                1120                1125

Gln Phe  Lys Lys Val Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe
    1130                1135                1140

Thr Gln  Pro Leu Tyr Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu
    1145                1150                1155

Leu Gly  Pro Tyr Ile Arg Ala  Glu Val Glu Asp Asn  Ile Met Val
    1160                1165                1170

Thr Phe  Arg Asn Gln Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser
    1175                1180                1185

Leu Ile  Ser Tyr Glu Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg
    1190                1195                1200

Lys Asn  Phe Val Lys Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys
    1205                1210                1215

Val Gln  His His Met Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys
    1220                1225                1230

Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu Glu Lys  Asp Val His
    1235                1240                1245

Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys His Thr  Asn Thr Leu
    1250                1255                1260

Asn Pro  Ala His Gly Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu
    1265                1270                1275

Phe Phe  Thr Ile Phe Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu
    1280                1285                1290

Asn Met  Glu Arg Asn Cys Arg  Gly Ala Pro Gly Thr  Pro Gly Ser
    1295                1300                1305

Gly Thr  Ala Ser Ser Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1310                1315                1320

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1325                1330                1335

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser
    1340                1345                1350

Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro Gly Ala  Ser Pro Gly
    1355                1360                1365
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1370 1375 1380

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
1385 1390 1395

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
1400 1405 1410

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
1415 1420 1425

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
1430 1435 1440

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
1445 1450 1455

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr
1460 1465 1470

Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
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Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
1490 1495 1500

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
1505 1510 1515

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser
1520 1525 1530

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser
1535 1540 1545

Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
1550 1555 1560

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
1565 1570 1575

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Ala Ser
1580 1585 1590

Ser Ala Pro Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu
1595 1600 1605

Asn Tyr Arg Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu
1610 1615 1620

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Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1625                1630                1635

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1640                1645                1650

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1655                1660                1665

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1670                1675                1680

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1685                1690                1695

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1700                1705                1710

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1715                1720                1725

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1730                1735                1740

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1745                1750                1755

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1760                1765                1770

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1775                1780                1785

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1790                1795                1800

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1805                1810                1815

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1820                1825                1830

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1835                1840                1845

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1850                1855                1860

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1865                1870                1875
```

Ile Ser Asp Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met
1880 1885 1890

Phe Ala Thr Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly
1895 1900 1905

Arg Ser Asn Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp
1910 1915 1920

Leu Gln Val Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr
1925 1930 1935

Thr Gln Gly Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu
1940 1945 1950

Phe Leu Ile Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe
1955 1960 1965

Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser
1970 1975 1980

Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg
1985 1990 1995

Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu
2000 2005 2010

Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala
2015 2020 2025

Gly Ser Pro Gly Ala Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu
2030 2035 2040

Glu Asp Leu Ser Pro Ala Thr Gly His His His His His His
2045 2050 2055

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atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

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atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

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<211> 1585
<212> PRT
<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic

polypeptide

<400> 1421

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
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Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
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530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
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Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Ser Pro Ala
740 745 750

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
755 760 765

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
770 775 780

Ala Ser Ser Pro Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg
785 790 795 800

Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile

805 810 815

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu
820 825 830

Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile
835 840 845

Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His
850 855 860

Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys
865 870 875 880

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
885 890 895

Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
900 905 910

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser
915 920 925

Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln
930 935 940

Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
945 950 955 960

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp
965 970 975

Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu
980 985 990

Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val Cys His Thr
995 1000 1005

Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr Val Gln Glu
1010 1015 1020

Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys Ser Trp Tyr
1025 1030 1035

Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Gly Ala Pro Gly Ser
1040 1045 1050

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
1055 1060 1065

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly

1070 1075 1080

Ser Glu Thr Pro Ala Ser Ser Ala Pro Cys Asn Ile Gln Met Glu
1085 1090 1095

Asp Pro Thr Phe Lys Glu Asn Tyr Arg Phe His Ala Ile Asn Gly
1100 1105 1110

Tyr Ile Met Asp Thr Leu Pro Gly Leu Val Met Ala Gln Asp Gln
1115 1120 1125

Arg Ile Arg Trp Tyr Leu Leu Ser Met Gly Ser Asn Glu Asn Ile
1130 1135 1140

His Ser Ile His Phe Ser Gly His Val Phe Thr Val Arg Lys Lys
1145 1150 1155

Glu Glu Tyr Lys Met Ala Leu Tyr Asn Leu Tyr Pro Gly Val Phe
1160 1165 1170

Glu Thr Val Glu Met Leu Pro Ser Lys Ala Gly Ile Trp Arg Val
1175 1180 1185

Glu Cys Leu Ile Gly Glu His Leu His Ala Gly Met Ser Thr Leu
1190 1195 1200

Phe Leu Val Tyr Ser Asn Lys Cys Gln Thr Pro Leu Gly Met Ala
1205 1210 1215

Ser Gly His Ile Arg Asp Phe Gln Ile Thr Ala Ser Gly Gln Tyr
1220 1225 1230

Gly Gln Trp Ala Pro Lys Leu Ala Arg Leu His Tyr Ser Gly Ser
1235 1240 1245

Ile Asn Ala Trp Ser Thr Lys Glu Pro Phe Ser Trp Ile Lys Val
1250 1255 1260

Asp Leu Leu Ala Pro Met Ile Ile His Gly Ile Lys Thr Gln Gly
1265 1270 1275

Ala Arg Gln Lys Phe Ser Ser Leu Tyr Ile Ser Gln Phe Ile Ile
1280 1285 1290

Met Tyr Ser Leu Asp Gly Lys Lys Trp Gln Thr Tyr Arg Gly Asn
1295 1300 1305

Ser Thr Gly Thr Leu Met Val Phe Phe Gly Asn Val Asp Ser Ser
1310 1315 1320

Gly Ile Lys His Asn Ile Phe Asn Pro Pro Ile Ile Ala Arg Tyr

1325 1330 1335

Ile Arg Leu His Pro Thr His Tyr Ser Ile Arg Ser Thr Leu Arg
1340 1345 1350

Met Glu Leu Met Gly Cys Asp Leu Asn Ser Cys Ser Met Pro Leu
1355 1360 1365

Gly Met Glu Ser Lys Ala Ile Ser Asp Ala Gln Ile Thr Ala Ser
1370 1375 1380

Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp Ser Pro Ser Lys Ala
1385 1390 1395

Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg Pro Gln Val
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Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln Lys Thr Met
1415 1420 1425

Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser Leu Leu Thr
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Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly
1445 1450 1455

His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val Lys Val Phe
1460 1465 1470

Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn Ser Leu Asp
1475 1480 1485

Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro Gln Ser Trp
1490 1495 1500

Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly Cys Glu Ala
1505 1510 1515

Gln Asp Leu Tyr Gly Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
1520 1525 1530

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1535 1540 1545

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Ala Ser
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Gly His His His His His His

1580 1585

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<211> 4758
<212> DNA
<213> Artificial Sequence

<220>
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<213> Artificial Sequence

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

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Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350
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Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620
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Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Ser Pro Ala
            740                 745                 750

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
        755                 760                 765

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
    770                 775                 780

Ala Ser Ser Pro Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg
785                 790                 795                 800

Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
                805                 810                 815

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu
            820                 825                 830

Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile
        835                 840                 845

Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His
    850                 855                 860

Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys
865                 870                 875                 880

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
                885                 890                 895
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Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
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Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser
        915                 920                 925

Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln
    930                 935                 940

Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
945                 950                 955                 960

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp
                965                 970                 975

Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu
            980                 985                 990

Lys Asp Val His Ser Gly Leu Ile  Gly Pro Leu Leu Val  Cys His Thr
        995                 1000                1005

Asn Thr  Leu Asn Pro Ala His  Gly Arg Gln Val Thr  Val Gln Glu
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Phe Ala  Leu Phe Phe Thr Ile  Phe Asp Glu Thr Lys  Ser Trp Tyr
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Phe Thr  Glu Asn Met Glu Arg  Asn Cys Arg Gly Ala  Pro Gly Thr
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Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro Gly Ala  Ser Pro Gly
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Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1070                1075                1080

Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    1085                1090                1095

Pro Gly  Ser Ser Pro Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Thr
    1100                1105                1110

Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro Gly Ala  Ser Pro Gly
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Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
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Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    1145                1150                1155
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Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser
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Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
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Thr Ser  Ser Thr Gly Ser Pro  Ala Ser Ser Ala Pro  Cys Asn Ile
    1190                1195                1200

Gln Met  Glu Asp Pro Thr Phe  Lys Glu Asn Tyr Arg  Phe His Ala
    1205                1210                1215

Ile Asn  Gly Tyr Ile Met Asp  Thr Leu Pro Gly Leu  Val Met Ala
    1220                1225                1230

Gln Asp  Gln Arg Ile Arg Trp  Tyr Leu Leu Ser Met  Gly Ser Asn
    1235                1240                1245

Glu Asn  Ile His Ser Ile His  Phe Ser Gly His Val  Phe Thr Val
    1250                1255                1260

Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
    1265                1270                1275

Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    1280                1285                1290

Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    1295                1300                1305

Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    1310                1315                1320

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
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Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
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Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
    1355                1360                1365

Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
    1370                1375                1380

Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
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Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
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Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
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Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
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Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1445                  1450                  1455

Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
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Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    1475                  1480                  1485

Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
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Ser Lys  Ala Arg Leu His Leu  Gln Gly Arg Ser Asn  Ala Trp Arg
    1505                  1510                  1515

Pro Gln  Val Asn Asn Pro Lys  Glu Trp Leu Gln Val  Asp Phe Gln
    1520                  1525                  1530

Lys Thr  Met Lys Val Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser
    1535                  1540                  1545

Leu Leu  Thr Ser Met Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser
    1550                  1555                  1560

Gln Asp  Gly His Gln Trp Thr  Leu Phe Phe Gln Asn  Gly Lys Val
    1565                  1570                  1575

Lys Val  Phe Gln Gly Asn Gln  Asp Ser Phe Thr Pro  Val Val Asn
    1580                  1585                  1590

Ser Leu  Asp Pro Pro Leu Leu  Thr Arg Tyr Leu Arg  Ile His Pro
    1595                  1600                  1605

Gln Ser  Trp Val His Gln Ile  Ala Leu Arg Met Glu  Val Leu Gly
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Cys Glu  Ala Gln Asp Leu Tyr  Gly Ala Pro Gly Ser  Pro Ala Gly
    1625                  1630                  1635

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
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Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
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Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
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Pro Ala Thr Gly His His His His His His
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gtgaataatc caaaagagtg gctgcaagtg gacttccaga agacaatgaa agtcacagga 4620

gtaactactc agggagtaaa atctctgctt accagcatgt atgtgaagga gttcctcatc 4680

tccagcagtc aagatggcca tcagtggact ctctttttc agaatggcaa agtaaaggtt 4740

tttcagggaa atcaagactc cttcacacct gtggtgaact ctctagaccc accgttactg 4800

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gttctgggct gcgaggcaca ggacctctac ggcgcgccag gttctcctgc tggctccccc 4920

acctcaacag aagaggggac aagcgaaagc gctacgcctg agagtggccc tggctctgag 4980

ccagccacct ccggctctga aacccctgcc tcgagcgaac aaaaactcat ctcagaagag 5040

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<210> 1425
<211> 1837
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1425
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro

```
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
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290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe

565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Ser Pro Ala
740 745 750

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
755 760 765

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
770 775 780

Ala Ser Ser Pro Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg
785 790 795 800

Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
805 810 815

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu
820 825 830

Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile

835 840 845

Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His
850 855 860

Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys
865 870 875 880

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
885 890 895

Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
900 905 910

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser
915 920 925

Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln
930 935 940

Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
945 950 955 960

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp
965 970 975

Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu
980 985 990

Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val Cys His Thr
995 1000 1005

Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr Val Gln Glu
1010 1015 1020

Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys Ser Trp Tyr
1025 1030 1035

Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Gly Ala Pro Gly Thr
1040 1045 1050

Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
1055 1060 1065

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
1070 1075 1080

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
1085 1090 1095

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala

1100 1105 1110

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
1115 1120 1125

Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
1130 1135 1140

Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
1145 1150 1155

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
1160 1165 1170

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
1175 1180 1185

Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
1190 1195 1200

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
1205 1210 1215

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser
1220 1225 1230

Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser
1235 1240 1245

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
1250 1255 1260

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
1265 1270 1275

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser
1280 1285 1290

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly
1295 1300 1305

Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
1310 1315 1320

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
1325 1330 1335

Pro Ala Ser Ser Ala Pro Cys Asn Ile Gln Met Glu Asp Pro Thr
1340 1345 1350

Phe Lys Glu Asn Tyr Arg Phe His Ala Ile Asn Gly Tyr Ile Met

1355 1360 1365

Asp Thr Leu Pro Gly Leu Val Met Ala Gln Asp Gln Arg Ile Arg
1370 1375 1380

Trp Tyr Leu Leu Ser Met Gly Ser Asn Glu Asn Ile His Ser Ile
1385 1390 1395

His Phe Ser Gly His Val Phe Thr Val Arg Lys Lys Glu Glu Tyr
1400 1405 1410

Lys Met Ala Leu Tyr Asn Leu Tyr Pro Gly Val Phe Glu Thr Val
1415 1420 1425

Glu Met Leu Pro Ser Lys Ala Gly Ile Trp Arg Val Glu Cys Leu
1430 1435 1440

Ile Gly Glu His Leu His Ala Gly Met Ser Thr Leu Phe Leu Val
1445 1450 1455

Tyr Ser Asn Lys Cys Gln Thr Pro Leu Gly Met Ala Ser Gly His
1460 1465 1470

Ile Arg Asp Phe Gln Ile Thr Ala Ser Gly Gln Tyr Gly Gln Trp
1475 1480 1485

Ala Pro Lys Leu Ala Arg Leu His Tyr Ser Gly Ser Ile Asn Ala
1490 1495 1500

Trp Ser Thr Lys Glu Pro Phe Ser Trp Ile Lys Val Asp Leu Leu
1505 1510 1515

Ala Pro Met Ile Ile His Gly Ile Lys Thr Gln Gly Ala Arg Gln
1520 1525 1530

Lys Phe Ser Ser Leu Tyr Ile Ser Gln Phe Ile Ile Met Tyr Ser
1535 1540 1545

Leu Asp Gly Lys Lys Trp Gln Thr Tyr Arg Gly Asn Ser Thr Gly
1550 1555 1560

Thr Leu Met Val Phe Phe Gly Asn Val Asp Ser Ser Gly Ile Lys
1565 1570 1575

His Asn Ile Phe Asn Pro Pro Ile Ile Ala Arg Tyr Ile Arg Leu
1580 1585 1590

His Pro Thr His Tyr Ser Ile Arg Ser Thr Leu Arg Met Glu Leu
1595 1600 1605

Met Gly Cys Asp Leu Asn Ser Cys Ser Met Pro Leu Gly Met Glu

1610 1615 1620

Ser Lys Ala Ile Ser Asp Ala Gln Ile Thr Ala Ser Ser Tyr Phe
1625 1630 1635

Thr Asn Met Phe Ala Thr Trp Ser Pro Ser Lys Ala Arg Leu His
1640 1645 1650

Leu Gln Gly Arg Ser Asn Ala Trp Arg Pro Gln Val Asn Asn Pro
1655 1660 1665

Lys Glu Trp Leu Gln Val Asp Phe Gln Lys Thr Met Lys Val Thr
1670 1675 1680

Gly Val Thr Thr Gln Gly Val Lys Ser Leu Leu Thr Ser Met Tyr
1685 1690 1695

Val Lys Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly His Gln Trp
1700 1705 1710

Thr Leu Phe Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn
1715 1720 1725

Gln Asp Ser Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu
1730 1735 1740

Leu Thr Arg Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln
1745 1750 1755

Ile Ala Leu Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu
1760 1765 1770

Tyr Gly Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
1775 1780 1785

Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
1790 1795 1800

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Ala Ser Ser Glu Gln
1805 1810 1815

Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala Thr Gly His His
1820 1825 1830

His His His His
1835

<210> 1426
<211> 5514
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

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gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

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gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

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cctggctctg agccagccac ctccggctct gaaacccctg cctcgagcga acaaaaactc 5460

atctcagaag aggatctgtc acctgcaacc ggtcatcatc accatcacca ttga 5514

<210> 1427
<211> 1693
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1427
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1 5 10 15

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285
```

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Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560
```

```
Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Thr Pro Gly
            740                 745                 750

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
        755                 760                 765

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    770                 775                 780

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
785                 790                 795                 800

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
                805                 810                 815

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            820                 825                 830
```

```
Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
        835                 840                 845

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
    850                 855                 860

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
865                 870                 875                 880

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095
```

```
Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1145                1150                1155

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr
    1160                1165                1170

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1175                1180                1185

Thr Ser  Gly Ser Glu Thr Pro  Ala Ser Ser Ala Pro  Cys Asn Ile
    1190                1195                1200

Gln Met  Glu Asp Pro Thr Phe  Lys Glu Asn Tyr Arg  Phe His Ala
    1205                1210                1215

Ile Asn  Gly Tyr Ile Met Asp  Thr Leu Pro Gly Leu  Val Met Ala
    1220                1225                1230

Gln Asp  Gln Arg Ile Arg Trp  Tyr Leu Leu Ser Met  Gly Ser Asn
    1235                1240                1245

Glu Asn  Ile His Ser Ile His  Phe Ser Gly His Val  Phe Thr Val
    1250                1255                1260

Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
    1265                1270                1275

Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    1280                1285                1290

Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    1295                1300                1305

Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    1310                1315                1320

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
    1325                1330                1335

Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
    1340                1345                1350
```

```
Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
    1355                1360                1365

Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
    1370                1375                1380

Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    1385                1390                1395

Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
    1400                1405                1410

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    1415                1420                1425

Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
    1430                1435                1440

Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1445                1450                1455

Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
    1460                1465                1470

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    1475                1480                1485

Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
    1490                1495                1500

Ser Lys  Ala Arg Leu His Leu  Gln Gly Arg Ser Asn  Ala Trp Arg
    1505                1510                1515

Pro Gln  Val Asn Asn Pro Lys  Glu Trp Leu Gln Val  Asp Phe Gln
    1520                1525                1530

Lys Thr  Met Lys Val Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser
    1535                1540                1545

Leu Leu  Thr Ser Met Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser
    1550                1555                1560

Gln Asp  Gly His Gln Trp Thr  Leu Phe Phe Gln Asn  Gly Lys Val
    1565                1570                1575

Lys Val  Phe Gln Gly Asn Gln  Asp Ser Phe Thr Pro  Val Val Asn
    1580                1585                1590

Ser Leu  Asp Pro Pro Leu Leu  Thr Arg Tyr Leu Arg  Ile His Pro
    1595                1600                1605
```

Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly
1610 1615 1620

Cys Glu Ala Gln Asp Leu Tyr Gly Ala Pro Gly Ser Pro Ala Gly
1625 1630 1635

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
1640 1645 1650

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1655 1660 1665

Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
1670 1675 1680

Pro Ala Thr Gly His His His His His His
1685 1690

<210> 1428
<211> 5082
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 1428
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ctcggtgagc tgcctgtgga cgcaagattt cctcctagag tgccaaaatc ttttccattc 120

aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

agagatctag cttcaggact cattggccct ctcctcatct gctacaaaga atctgtagat 1680

caaagaggaa accagataat gtcagacaag aggaatgtca tcctgttttc tgtatttgat 1740

gagaaccgaa gctggtacct cacagagaat atacaacgct ttctccccaa tccagctgga 1800

gtgcagcttg aggatccaga gttccaagcc tccaacatca tgcacagcat caatggctat 1860

gtttttgata gtttgcagtt gtcagtttgt ttgcatgagg tggcatactg gtacattcta 1920

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gacagttatg aagatatttc agcatacttg ctgagtaaaa acaatgccat tgaaccaaga 2220

agcttctctc aaaacggcgc gccaggtaca ccgggcagcg gcacggcttc gtcgtcaccc 2280

ggctcgtcca caccgtcggg agctacggga agcccaggag cgtcaccggg aacgtcgtca 2340

acggggtcac cgggtacgcc aggtagcggc acggccagca gctcgccagg ttcatcgacc 2400

ccgtcgggag cgactgggtc gcccggatca agcccgtcag cttccactgg aacaggaccc 2460

gggtcgtcgc cgtcagcctc aacggggaca ggacctggtt catcgacgcc gtcaggggcg 2520

acaggctcgc ccggatcgtc aacaccctcg ggggcaacgg ggagccctgg tgcgtcgcct 2580

ggaacctcat ccaccggaag cccgggggcc tcgccgggta cgagctccac gggatcgccc 2640

ggagcgtccc ccggaacttc aagcacaggg agccctgcct cgagcccacc agtcttgaaa 2700

cgccatcaag ctgaaataac tcgtactact cttcagtcag atcaagagga aatcgattat 2760

gatgatacca tatcagttga aatgaagaag gaagattttg acatttatga tgaggatgaa 2820

aatcagagcc cccgcagctt tcaaaagaaa acacgacact attttattgc tgcagtggag 2880

aggctctggg attatgggat gagtagctcc ccacatgttc taagaaacag ggctcagagt 2940

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cagcccttat accgtggaga actaaatgaa catttgggac tcctggggcc atatataaga 3060

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cctgctcatg ggagacaagt gacagtacag gaatttgctc tgtttttcac catctttgat 3420

gagaccaaaa gctggtactt cactgaaaat atggaaagaa actgcagggg cgcgccaggt 3480

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agtggccctg gctctgagcc agccacctcc ggctctgaaa cccctgcctc gagcgctccc 3600

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agcaccaagg agccctttc ttggatcaag gtggatctgt tggcaccaat gattattcac 4140

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atcatgtata gtcttgatgg gaagaagtgg cagacttatc gaggaaattc cactggaacc 4260

ttaatggtct tctttggcaa tgtggattca tctgggataa aacacaatat ttttaaccct 4320

ccaattattg ctcgatacat ccgtttgcac ccaactcatt atagcattcg cagcactctt 4380

cgcatggagt tgatgggctg tgatttaaat agttgcagca tgccattggg aatggagagt 4440

aaagcaatat cagatgcaca gattactgct tcatcctact ttaccaatat gtttgccacc 4500

tggtctcctt caaaagctcg acttcacctc caagggagga gtaatgcctg gagacctcag 4560

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tccagcagtc aagatggcca tcagtggact ctctttttc agaatggcaa agtaaaggtt 4740

tttcagggaa atcaagactc cttcacacct gtggtgaact ctctagaccc accgttactg 4800

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acctcaacag aagaggggac aagcgaaagc gctacgcctg agagtggccc tggctctgag 4980

ccagccacct ccggctctga aacccctgcc tcgagcgaac aaaaactcat ctcagaagag 5040

gatctgtcac ctgcaaccgg tcatcatcac catcaccatt ga 5082

<210> 1429
<211> 1801
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1429

```
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1               5                   10                  15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
```

225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys

```
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Thr Pro Gly
            740                 745                 750

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
        755                 760                 765

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
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770 775 780

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
785 790 795 800

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
805 810 815

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
820 825 830

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
835 840 845

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
850 855 860

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
865 870 875 880

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Pro
885 890 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
900 905 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
915 920 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
930 935 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945 950 955 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
965 970 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
980 985 990

Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
995 1000 1005

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1010 1015 1020

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1025 1030 1035

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg

```
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1145                1150                1155

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala
    1160                1165                1170

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1175                1180                1185

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1190                1195                1200

Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
    1205                1210                1215

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala
    1220                1225                1230

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1235                1240                1245

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1250                1255                1260

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1265                1270                1275

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ala
    1280                1285                1290

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Ala Ser  Ser Ala Pro
```

1295 1300 1305

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1310 1315 1320

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1325 1330 1335

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1340 1345 1350

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1355 1360 1365

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1370 1375 1380

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1385 1390 1395

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1400 1405 1410

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1415 1420 1425

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1430 1435 1440

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1445 1450 1455

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1460 1465 1470

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1475 1480 1485

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1490 1495 1500

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1505 1510 1515

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1520 1525 1530

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1535 1540 1545

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser

1550 1555 1560

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1565 1570 1575

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1580 1585 1590

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1595 1600 1605

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1610 1615 1620

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1625 1630 1635

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1640 1645 1650

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1655 1660 1665

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1670 1675 1680

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1685 1690 1695

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1700 1705 1710

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1715 1720 1725

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Ala Pro Gly Ser
1730 1735 1740

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
1745 1750 1755

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
1760 1765 1770

Ser Glu Thr Pro Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu
1775 1780 1785

Asp Leu Ser Pro Ala Thr Gly His His His His His His
1790 1795 1800

<210> 1430

<211> 5406
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

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gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

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cctgagagtg gccctggctc tgagccagcc acctccggct ctgaaacccc tgcctcgagc 5340

gaacaaaaac tcatctcaga agaggatctg tcacctgcaa ccggtcatca tcaccatcac 5400

cattga 5406

<210> 1431
<211> 1945
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1431
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285
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Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560
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Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Thr Pro Gly
            740                 745                 750

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
        755                 760                 765

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    770                 775                 780

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
785                 790                 795                 800

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
                805                 810                 815

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            820                 825                 830
```

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Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
        835                 840                 845

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
    850                 855                 860

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
865                 870                 875                 880

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095
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```
Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1145                1150                1155

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser
    1160                1165                1170

Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Ser  Ser Thr Pro
    1175                1180                1185

Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1190                1195                1200

Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
    1205                1210                1215

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1220                1225                1230

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Thr  Pro Gly Ser
    1235                1240                1245

Gly Thr  Ala Ser Ser Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1250                1255                1260

Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    1265                1270                1275

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1280                1285                1290

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Thr  Pro Gly Ser
    1295                1300                1305

Gly Thr  Ala Ser Ser Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1310                1315                1320

Thr Gly  Ser Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1325                1330                1335

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Thr
    1340                1345                1350
```

```
Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro
    1355                1360                1365

Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1370                1375                1380

Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
    1385                1390                1395

Pro Gly  Ser Ser Pro Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ala
    1400                1405                1410

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Thr  Pro Gly Ser
    1415                1420                1425

Gly Thr  Ala Ser Ser Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1430                1435                1440

Thr Gly  Ser Pro Ala Ser Ser  Ala Pro Cys Asn Ile  Gln Met Glu
    1445                1450                1455

Asp Pro  Thr Phe Lys Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly
    1460                1465                1470

Tyr Ile  Met Asp Thr Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln
    1475                1480                1485

Arg Ile  Arg Trp Tyr Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile
    1490                1495                1500

His Ser  Ile His Phe Ser Gly  His Val Phe Thr Val  Arg Lys Lys
    1505                1510                1515

Glu Glu  Tyr Lys Met Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe
    1520                1525                1530

Glu Thr  Val Glu Met Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val
    1535                1540                1545

Glu Cys  Leu Ile Gly Glu His  Leu His Ala Gly Met  Ser Thr Leu
    1550                1555                1560

Phe Leu  Val Tyr Ser Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala
    1565                1570                1575

Ser Gly  His Ile Arg Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr
    1580                1585                1590

Gly Gln  Trp Ala Pro Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser
    1595                1600                1605
```

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Ile Asn  Ala Trp Ser Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val
    1610                 1615                 1620

Asp Leu  Leu Ala Pro Met Ile  Ile His Gly Ile Lys  Thr Gln Gly
    1625                 1630                 1635

Ala Arg  Gln Lys Phe Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile
    1640                 1645                 1650

Met Tyr  Ser Leu Asp Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn
    1655                 1660                 1665

Ser Thr  Gly Thr Leu Met Val  Phe Phe Gly Asn Val  Asp Ser Ser
    1670                 1675                 1680

Gly Ile  Lys His Asn Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr
    1685                 1690                 1695

Ile Arg  Leu His Pro Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg
    1700                 1705                 1710

Met Glu  Leu Met Gly Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu
    1715                 1720                 1725

Gly Met  Glu Ser Lys Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser
    1730                 1735                 1740

Ser Tyr  Phe Thr Asn Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala
    1745                 1750                 1755

Arg Leu  His Leu Gln Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val
    1760                 1765                 1770

Asn Asn  Pro Lys Glu Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met
    1775                 1780                 1785

Lys Val  Thr Gly Val Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr
    1790                 1795                 1800

Ser Met  Tyr Val Lys Glu Phe  Leu Ile Ser Ser Ser  Gln Asp Gly
    1805                 1810                 1815

His Gln  Trp Thr Leu Phe Phe  Gln Asn Gly Lys Val  Lys Val Phe
    1820                 1825                 1830

Gln Gly  Asn Gln Asp Ser Phe  Thr Pro Val Val Asn  Ser Leu Asp
    1835                 1840                 1845

Pro Pro  Leu Leu Thr Arg Tyr  Leu Arg Ile His Pro  Gln Ser Trp
    1850                 1855                 1860
```

Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly Cys Glu Ala
1865 1870 1875

Gln Asp Leu Tyr Gly Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
1880 1885 1890

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1895 1900 1905

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Ala Ser
1910 1915 1920

Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala Thr
1925 1930 1935

Gly His His His His His His
1940 1945

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<211> 5838
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

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caaagaggaa accagataat gtcagacaag aggaatgtca tcctgttttc tgtatttgat 1740

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catcaccatc accattga 5838

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<211> 1837
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val

```
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
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385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro

660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Pro Gly Ala Ser
740 745 750

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
755 760 765

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
770 775 780

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro
785 790 795 800

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
805 810 815

Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
820 825 830

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
835 840 845

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
850 855 860

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
865 870 875 880

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
885 890 895

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
900 905 910

Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
915 920 925

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser

930 935 940

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
945 950 955 960

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
965 970 975

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
980 985 990

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
995 1000 1005

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
1010 1015 1020

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Ala Ser
1025 1030 1035

Ser Pro Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr
1040 1045 1050

Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
1055 1060 1065

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp
1070 1075 1080

Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr
1085 1090 1095

Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser
1100 1105 1110

Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro
1115 1120 1125

Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe
1130 1135 1140

Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
1145 1150 1155

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val
1160 1165 1170

Thr Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser
1175 1180 1185

Leu Ile Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg

1190 1195 1200

Lys Asn Phe Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys
1205 1210 1215

Val Gln His His Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys
1220 1225 1230

Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu Lys Asp Val His
1235 1240 1245

Ser Gly Leu Ile Gly Pro Leu Leu Val Cys His Thr Asn Thr Leu
1250 1255 1260

Asn Pro Ala His Gly Arg Gln Val Thr Val Gln Glu Phe Ala Leu
1265 1270 1275

Phe Phe Thr Ile Phe Asp Glu Thr Lys Ser Trp Tyr Phe Thr Glu
1280 1285 1290

Asn Met Glu Arg Asn Cys Arg Gly Ala Pro Gly Ser Pro Ala Gly
1295 1300 1305

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
1310 1315 1320

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1325 1330 1335

Pro Ala Ser Ser Ala Pro Cys Asn Ile Gln Met Glu Asp Pro Thr
1340 1345 1350

Phe Lys Glu Asn Tyr Arg Phe His Ala Ile Asn Gly Tyr Ile Met
1355 1360 1365

Asp Thr Leu Pro Gly Leu Val Met Ala Gln Asp Gln Arg Ile Arg
1370 1375 1380

Trp Tyr Leu Leu Ser Met Gly Ser Asn Glu Asn Ile His Ser Ile
1385 1390 1395

His Phe Ser Gly His Val Phe Thr Val Arg Lys Lys Glu Glu Tyr
1400 1405 1410

Lys Met Ala Leu Tyr Asn Leu Tyr Pro Gly Val Phe Glu Thr Val
1415 1420 1425

Glu Met Leu Pro Ser Lys Ala Gly Ile Trp Arg Val Glu Cys Leu
1430 1435 1440

Ile Gly Glu His Leu His Ala Gly Met Ser Thr Leu Phe Leu Val

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Tyr Ser  Asn Lys Cys Gln Thr  Pro Leu Gly Met Ala  Ser Gly His
    1460                1465                1470

Ile Arg  Asp Phe Gln Ile Thr  Ala Ser Gly Gln Tyr  Gly Gln Trp
    1475                1480                1485

Ala Pro  Lys Leu Ala Arg Leu  His Tyr Ser Gly Ser  Ile Asn Ala
    1490                1495                1500

Trp Ser  Thr Lys Glu Pro Phe  Ser Trp Ile Lys Val  Asp Leu Leu
    1505                1510                1515

Ala Pro  Met Ile Ile His Gly  Ile Lys Thr Gln Gly  Ala Arg Gln
    1520                1525                1530

Lys Phe  Ser Ser Leu Tyr Ile  Ser Gln Phe Ile Ile  Met Tyr Ser
    1535                1540                1545

Leu Asp  Gly Lys Lys Trp Gln  Thr Tyr Arg Gly Asn  Ser Thr Gly
    1550                1555                1560

Thr Leu  Met Val Phe Phe Gly  Asn Val Asp Ser Ser  Gly Ile Lys
    1565                1570                1575

His Asn  Ile Phe Asn Pro Pro  Ile Ile Ala Arg Tyr  Ile Arg Leu
    1580                1585                1590

His Pro  Thr His Tyr Ser Ile  Arg Ser Thr Leu Arg  Met Glu Leu
    1595                1600                1605

Met Gly  Cys Asp Leu Asn Ser  Cys Ser Met Pro Leu  Gly Met Glu
    1610                1615                1620

Ser Lys  Ala Ile Ser Asp Ala  Gln Ile Thr Ala Ser  Ser Tyr Phe
    1625                1630                1635

Thr Asn  Met Phe Ala Thr Trp  Ser Pro Ser Lys Ala  Arg Leu His
    1640                1645                1650

Leu Gln  Gly Arg Ser Asn Ala  Trp Arg Pro Gln Val  Asn Asn Pro
    1655                1660                1665

Lys Glu  Trp Leu Gln Val Asp  Phe Gln Lys Thr Met  Lys Val Thr
    1670                1675                1680

Gly Val  Thr Thr Gln Gly Val  Lys Ser Leu Leu Thr  Ser Met Tyr
    1685                1690                1695

Val Lys  Glu Phe Leu Ile Ser  Ser Ser Gln Asp Gly  His Gln Trp
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1700 1705 1710

Thr Leu Phe Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn
1715 1720 1725

Gln Asp Ser Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu
1730 1735 1740

Leu Thr Arg Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln
1745 1750 1755

Ile Ala Leu Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu
1760 1765 1770

Tyr Gly Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
1775 1780 1785

Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
1790 1795 1800

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Ala Ser Ser Glu Gln
1805 1810 1815

Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala Thr Gly His His
1820 1825 1830

His His His His
1835

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<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic polynucleotide

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gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

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ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

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Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
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Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380
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Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655
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Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Pro Gly Ala Ser
            740                 745                 750

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
        755                 760                 765

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
    770                 775                 780

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro
785                 790                 795                 800

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
                805                 810                 815

Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
            820                 825                 830

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
        835                 840                 845

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
    850                 855                 860

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
865                 870                 875                 880

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
                885                 890                 895

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
            900                 905                 910

Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
        915                 920                 925
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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
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Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
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Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
            980                 985                 990

Pro Gly Thr Ser Ser Thr Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser
        995                 1000                1005

Thr Gly  Thr Gly Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
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Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Ala Ser
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Ser Pro  Pro Val Leu Lys Arg  His Gln Ala Glu Ile  Thr Arg Thr
    1040                1045                1050

Thr Leu  Gln Ser Asp Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile
    1055                1060                1065

Ser Val  Glu Met Lys Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp
    1070                1075                1080

Glu Asn  Gln Ser Pro Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr
    1085                1090                1095

Phe Ile  Ala Ala Val Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser
    1100                1105                1110

Ser Pro  His Val Leu Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro
    1115                1120                1125

Gln Phe  Lys Lys Val Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe
    1130                1135                1140

Thr Gln  Pro Leu Tyr Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu
    1145                1150                1155

Leu Gly  Pro Tyr Ile Arg Ala  Glu Val Glu Asp Asn  Ile Met Val
    1160                1165                1170

Thr Phe  Arg Asn Gln Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser
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Leu Ile  Ser Tyr Glu Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg
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Lys Asn  Phe Val Lys Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys
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Val Gln  His His Met Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys
    1220                 1225                 1230

Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu Glu Lys  Asp Val His
    1235                 1240                 1245

Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys His Thr  Asn Thr Leu
    1250                 1255                 1260

Asn Pro  Ala His Gly Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu
    1265                 1270                 1275

Phe Phe  Thr Ile Phe Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu
    1280                 1285                 1290

Asn Met  Glu Arg Asn Cys Arg  Gly Ala Pro Gly Thr  Pro Gly Ser
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Gly Thr  Ala Ser Ser Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
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Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
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Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ser
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Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro Gly Thr  Pro Gly Ser
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Gly Thr  Ala Ser Ser Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1370                 1375                 1380

Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    1385                 1390                 1395

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ser
    1400                 1405                 1410

Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Ser  Ser Thr Pro
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Ser Gly  Ala Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
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Thr Gly  Ser Pro Ala Ser Ser  Ala Pro Cys Asn Ile  Gln Met Glu
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Asp Pro  Thr Phe Lys Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly
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Tyr Ile  Met Asp Thr Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln
    1475                1480                1485

Arg Ile  Arg Trp Tyr Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile
    1490                1495                1500

His Ser  Ile His Phe Ser Gly  His Val Phe Thr Val  Arg Lys Lys
    1505                1510                1515

Glu Glu  Tyr Lys Met Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe
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Glu Thr  Val Glu Met Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val
    1535                1540                1545

Glu Cys  Leu Ile Gly Glu His  Leu His Ala Gly Met  Ser Thr Leu
    1550                1555                1560

Phe Leu  Val Tyr Ser Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala
    1565                1570                1575

Ser Gly  His Ile Arg Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr
    1580                1585                1590

Gly Gln  Trp Ala Pro Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser
    1595                1600                1605

Ile Asn  Ala Trp Ser Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val
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Asp Leu  Leu Ala Pro Met Ile  Ile His Gly Ile Lys  Thr Gln Gly
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Ala Arg  Gln Lys Phe Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile
    1640                1645                1650

Met Tyr  Ser Leu Asp Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn
    1655                1660                1665

Ser Thr  Gly Thr Leu Met Val  Phe Phe Gly Asn Val  Asp Ser Ser
    1670                1675                1680

Gly Ile  Lys His Asn Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr
    1685                1690                1695
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Ile Arg  Leu His Pro Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg
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Gly Met  Glu Ser Lys Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser
    1730                1735                1740

Ser Tyr  Phe Thr Asn Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala
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Arg Leu  His Leu Gln Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val
    1760                1765                1770

Asn Asn  Pro Lys Glu Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met
    1775                1780                1785

Lys Val  Thr Gly Val Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr
    1790                1795                1800

Ser Met  Tyr Val Lys Glu Phe  Leu Ile Ser Ser Ser  Gln Asp Gly
    1805                1810                1815

His Gln  Trp Thr Leu Phe Phe  Gln Asn Gly Lys Val  Lys Val Phe
    1820                1825                1830

Gln Gly  Asn Gln Asp Ser Phe  Thr Pro Val Val Asn  Ser Leu Asp
    1835                1840                1845

Pro Pro  Leu Leu Thr Arg Tyr  Leu Arg Ile His Pro  Gln Ser Trp
    1850                1855                1860

Val His  Gln Ile Ala Leu Arg  Met Glu Val Leu Gly  Cys Glu Ala
    1865                1870                1875

Gln Asp  Leu Tyr Gly Ala Pro  Gly Ser Pro Ala Gly  Ser Pro Thr
    1880                1885                1890

Ser Thr  Glu Glu Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1895                1900                1905

Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser Glu Thr  Pro Ala Ser
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Ser Glu  Gln Lys Leu Ile Ser  Glu Glu Asp Leu Ser  Pro Ala Thr
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Gly His  His His His His His
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20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

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Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495
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His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Pro Gly Ala Ser
            740                 745                 750

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
        755                 760                 765
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Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
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Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro
785                 790                 795                 800

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
                805                 810                 815

Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
            820                 825                 830

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
        835                 840                 845

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
    850                 855                 860

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
865                 870                 875                 880

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
                885                 890                 895

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
            900                 905                 910

Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
        915                 920                 925

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
    930                 935                 940

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
945                 950                 955                 960

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
                965                 970                 975

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
            980                 985                 990

Pro Gly Thr Ser Ser Thr Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser
        995                 1000                1005

Thr Gly  Thr Gly Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
    1010                1015                1020

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Ala Ser
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Ser Pro  Pro Val Leu Lys Arg  His Gln Ala Glu Ile  Thr Arg Thr
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Thr Leu  Gln Ser Asp Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile
    1055                1060                1065

Ser Val  Glu Met Lys Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp
    1070                1075                1080

Glu Asn  Gln Ser Pro Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr
    1085                1090                1095

Phe Ile  Ala Ala Val Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser
    1100                1105                1110

Ser Pro  His Val Leu Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro
    1115                1120                1125

Gln Phe  Lys Lys Val Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe
    1130                1135                1140

Thr Gln  Pro Leu Tyr Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu
    1145                1150                1155

Leu Gly  Pro Tyr Ile Arg Ala  Glu Val Glu Asp Asn  Ile Met Val
    1160                1165                1170

Thr Phe  Arg Asn Gln Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser
    1175                1180                1185

Leu Ile  Ser Tyr Glu Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg
    1190                1195                1200

Lys Asn  Phe Val Lys Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys
    1205                1210                1215

Val Gln  His His Met Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys
    1220                1225                1230

Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu Glu Lys  Asp Val His
    1235                1240                1245

Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys His Thr  Asn Thr Leu
    1250                1255                1260

Asn Pro  Ala His Gly Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu
    1265                1270                1275

Phe Phe  Thr Ile Phe Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu
    1280                1285                1290
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Asn Met  Glu Arg Asn Cys Arg  Gly Ala Pro Gly Thr  Pro Gly Ser
    1295                 1300                 1305

Gly Thr  Ala Ser Ser Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
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Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1325                 1330                 1335

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser
    1340                 1345                 1350

Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro Gly Ala  Ser Pro Gly
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Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1370                 1375                 1380

Thr Gly  Ser Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1385                 1390                 1395

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1400                 1405                 1410

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1415                 1420                 1425

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1430                 1435                 1440

Thr Gly  Ser Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1445                 1450                 1455

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Thr
    1460                 1465                 1470

Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro
    1475                 1480                 1485

Ser Gly  Ala Thr Gly Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala
    1490                 1495                 1500

Ser Ser  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1505                 1510                 1515

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser
    1520                 1525                 1530

Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro Gly Ser  Ser Pro Ser
    1535                 1540                 1545
```

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Ala Ser  Thr Gly Thr Gly Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
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Thr Gly  Ser Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
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Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Ala Ser
    1580                1585                1590

Ser Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1595                1600                1605

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1610                1615                1620

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1625                1630                1635

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1640                1645                1650

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1655                1660                1665

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1670                1675                1680

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1685                1690                1695

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1700                1705                1710

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1715                1720                1725

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1730                1735                1740

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1745                1750                1755

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1760                1765                1770

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1775                1780                1785

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1790                1795                1800
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Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1805                1810                1815

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1820                1825                1830

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1835                1840                1845

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1850                1855                1860

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1865                1870                1875

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1880                1885                1890

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1895                1900                1905

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1910                1915                1920

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1925                1930                1935

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1940                1945                1950

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1955                1960                1965

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1970                1975                1980

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    1985                1990                1995

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    2000                2005                2010

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    2015                2020                2025

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr
    2030                2035                2040

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    2045                2050                2055
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Thr Ser Gly Ser Glu Thr Pro Ala Ser Ser Glu Gln Lys Leu Ile
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Ser Glu Glu Asp Leu Ser Pro Ala Thr Gly His His His His His
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His

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

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Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
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Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380
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Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
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Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655
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Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Ser Pro Ala
            740                 745                 750

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
        755                 760                 765

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
    770                 775                 780

Ala Ser Ser Pro Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg
785                 790                 795                 800

Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
                805                 810                 815

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu
            820                 825                 830

Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile
        835                 840                 845

Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His
    850                 855                 860

Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys
865                 870                 875                 880

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
                885                 890                 895

Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
            900                 905                 910

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser
        915                 920                 925
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Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln
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Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
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Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp
                965                 970                 975

Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu
            980                 985                 990

Lys Asp Val His Ser Gly Leu Ile  Gly Pro Leu Leu Val  Cys His Thr
        995                 1000                1005

Asn Thr  Leu Asn Pro Ala His  Gly Arg Gln Val Thr  Val Gln Glu
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Phe Ala  Leu Phe Phe Thr Ile  Phe Asp Glu Thr Lys  Ser Trp Tyr
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Phe Thr  Glu Asn Met Glu Arg  Asn Cys Arg Gly Ala  Pro Gly Ser
    1040                1045                1050

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    1055                1060                1065

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1070                1075                1080

Ser Glu  Thr Pro Ala Ser Ser  Ala Pro Cys Asn Ile  Gln Met Glu
    1085                1090                1095

Asp Pro  Thr Phe Lys Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly
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Tyr Ile  Met Asp Thr Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln
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Arg Ile  Arg Trp Tyr Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile
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His Ser  Ile His Phe Ser Gly  His Val Phe Thr Val  Arg Lys Lys
    1145                1150                1155

Glu Glu  Tyr Lys Met Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe
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Glu Thr  Val Glu Met Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val
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Glu Cys  Leu Ile Gly Glu His  Leu His Ala Gly Met  Ser Thr Leu
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Phe Leu  Val Tyr Ser Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala
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Ser Gly  His Ile Arg Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr
    1220                 1225                 1230

Gly Gln  Trp Ala Pro Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser
    1235                 1240                 1245

Ile Asn  Ala Trp Ser Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val
    1250                 1255                 1260

Asp Leu  Leu Ala Pro Met Ile  Ile His Gly Ile Lys  Thr Gln Gly
    1265                 1270                 1275

Ala Arg  Gln Lys Phe Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile
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Met Tyr  Ser Leu Asp Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn
    1295                 1300                 1305

Ser Thr  Gly Thr Leu Met Val  Phe Phe Gly Asn Val  Asp Ser Ser
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Gly Ile  Lys His Asn Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr
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Ile Arg  Leu His Pro Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg
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Met Glu  Leu Met Gly Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu
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Gly Met  Glu Ser Lys Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser
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Ser Tyr  Phe Thr Asn Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala
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Arg Leu  His Leu Gln Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val
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Asn Asn  Pro Lys Glu Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met
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Lys Val  Thr Gly Val Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr
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Ser Met  Tyr Val Lys Glu Phe  Leu Ile Ser Ser Ser  Gln Asp Gly
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His Gln  Trp Thr Leu Phe Phe  Gln Asn Gly Lys Val  Lys Val Phe
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Gln Gly  Asn Gln Asp Ser Phe  Thr Pro Val Val Asn  Ser Leu Asp
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Pro Pro  Leu Leu Thr Arg Tyr  Leu Arg Ile His Pro  Gln Ser Trp
    1490                1495                1500

Val His  Gln Ile Ala Leu Arg  Met Glu Val Leu Gly  Cys Glu Ala
    1505                1510                1515

Gln Asp  Leu Tyr Gly Ala Pro  Pro Gly Ser Ser Pro  Ser Ala Ser
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Thr Gly  Thr Gly Pro Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr
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Gly Pro  Gly Thr Pro Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly
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Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Pro
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Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ala Ser Pro  Gly Thr Ser
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Ser Thr  Gly Ser Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
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Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly
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Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser Pro Gly  Ala Ser Pro
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Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
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Ser Thr  Gly Ser Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
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Ser Ala  Ser Ser Glu Gln Lys  Leu Ile Ser Glu Glu  Asp Leu Ser
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Pro Ala  Thr Gly His His His  His His His
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aagtggcaga cttatcgagg aaattccact ggaaccttaa tggtcttctt tggcaatgtg 3960

gattcatctg ggataaaaca caatattttt aaccctccaa ttattgctcg atacatccgt 4020

ttgcacccaa ctcattatag cattcgcagc actcttcgca tggagttgat gggctgtgat 4080

ttaaatagtt gcagcatgcc attgggaatg gagagtaaag caatatcaga tgcacagatt 4140

actgcttcat cctactttac caatatgttt gccacctggt ctccttcaaa agctcgactt 4200

cacctccaag ggaggagtaa tgcctggaga cctcaggtga ataatccaaa agagtggctg 4260

caagtggact tccagaagac aatgaaagtc acaggagtaa ctactcaggg agtaaaatct 4320

ctgcttacca gcatgtatgt gaaggagttc ctcatctcca gcagtcaaga tggccatcag 4380

tggactctct tttttcagaa tggcaaagta aaggtttttc agggaaatca agactccttc 4440

acacctgtgg tgaactctct agacccaccg ttactgactc gctaccttcg aattcacccc 4500

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tcgcctggtt cgtccacccc gtcaggagcc acgggttcgc cgggaagcag cccaagcgca 4740

tccactggta cagggcctgg ggcttcaccg ggtacttcat ccacggggtc accgggaacg 4800

cccggatcgg ggacggcttc ctcatcacca ggatcgtcaa caccctcggg cgcaacgggc 4860

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agctcgacag gatcgcctgg ggcgtcaccc ggcacgtcga gcacaggcag ccccggaacc 4980

cctggatcgg gaaccgcgtc gtcaagcgcc tcgagcgaac aaaaactcat ctcagaagag 5040

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<210> 1441
<211> 1801
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1441
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro

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    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
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                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
```

595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Ser Pro Ala
740 745 750

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
755 760 765

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
770 775 780

Ala Ser Ser Pro Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg
785 790 795 800

Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
805 810 815

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu
820 825 830

Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile
835 840 845

Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His
850 855 860

Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys

865 870 875 880

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
885 890 895

Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
900 905 910

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser
915 920 925

Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln
930 935 940

Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
945 950 955 960

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp
965 970 975

Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu
980 985 990

Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val Cys His Thr
995 1000 1005

Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr Val Gln Glu
1010 1015 1020

Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys Ser Trp Tyr
1025 1030 1035

Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Gly Ala Pro Gly Thr
1040 1045 1050

Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly
1055 1060 1065

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
1070 1075 1080

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
1085 1090 1095

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr
1100 1105 1110

Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly
1115 1120 1125

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser

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    1130                1135                1140

Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    1145                1150                1155

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser
    1160                1165                1170

Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1175                1180                1185

Thr Ser  Ser Thr Gly Ser Pro  Ala Ser Ser Ala Pro  Cys Asn Ile
    1190                1195                1200

Gln Met  Glu Asp Pro Thr Phe  Lys Glu Asn Tyr Arg  Phe His Ala
    1205                1210                1215

Ile Asn  Gly Tyr Ile Met Asp  Thr Leu Pro Gly Leu  Val Met Ala
    1220                1225                1230

Gln Asp  Gln Arg Ile Arg Trp  Tyr Leu Leu Ser Met  Gly Ser Asn
    1235                1240                1245

Glu Asn  Ile His Ser Ile His  Phe Ser Gly His Val  Phe Thr Val
    1250                1255                1260

Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
    1265                1270                1275

Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    1280                1285                1290

Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    1295                1300                1305

Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    1310                1315                1320

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
    1325                1330                1335

Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
    1340                1345                1350

Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
    1355                1360                1365

Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
    1370                1375                1380

Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
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1385 1390 1395

Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp Gln Thr Tyr
1400 1405 1410

Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe Gly Asn Val
1415 1420 1425

Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro Pro Ile Ile
1430 1435 1440

Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser Ile Arg Ser
1445 1450 1455

Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn Ser Cys Ser
1460 1465 1470

Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp Ala Gln Ile
1475 1480 1485

Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp Ser Pro
1490 1495 1500

Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg
1505 1510 1515

Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln
1520 1525 1530

Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser
1535 1540 1545

Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser
1550 1555 1560

Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val
1565 1570 1575

Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn
1580 1585 1590

Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro
1595 1600 1605

Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly
1610 1615 1620

Cys Glu Ala Gln Asp Leu Tyr Gly Ala Pro Pro Gly Ser Ser Pro
1625 1630 1635

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser

1640 1645 1650

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
1655 1660 1665

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
1670 1675 1680

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
1685 1690 1695

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
1700 1705 1710

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
1715 1720 1725

Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
1730 1735 1740

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
1745 1750 1755

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
1760 1765 1770

Ala Ser Ser Ser Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu
1775 1780 1785

Asp Leu Ser Pro Ala Thr Gly His His His His His His
1790 1795 1800

<210> 1442
<211> 5406
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540
gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600
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gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720
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tcaacaccct cgggcgcaac gggcagcccc ggaacccctg gttcgggtac ggcgtcgtcg 5220

agccccggtg cgagcccggg aacaagctcg acaggatcgc ctggggcgtc acccggcacg 5280

tcgagcacag gcagccccgg aacccctgga tcgggaaccg cgtcgtcaag cgcctcgagc 5340

gaacaaaaac tcatctcaga agaggatctg tcacctgcaa ccggtcatca tcaccatcac 5400

cattga 5406

<210> 1443
<211> 1945
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

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Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

```
Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380
```

```
Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655
```

```
Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Ser Pro Ala
            740                 745                 750

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
        755                 760                 765

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
    770                 775                 780

Ala Ser Ser Pro Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg
785                 790                 795                 800

Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
                805                 810                 815

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu
            820                 825                 830

Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile
        835                 840                 845

Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His
    850                 855                 860

Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys
865                 870                 875                 880

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
                885                 890                 895

Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
            900                 905                 910

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser
        915                 920                 925
```

```
Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln
    930                 935                 940

Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
945                 950                 955                 960

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp
                965                 970                 975

Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu
            980                 985                 990

Lys Asp Val His Ser Gly Leu Ile  Gly Pro Leu Leu Val  Cys His Thr
        995                 1000                1005

Asn Thr  Leu Asn Pro Ala His  Gly Arg Gln Val Thr  Val Gln Glu
    1010                1015                1020

Phe Ala  Leu Phe Phe Thr Ile  Phe Asp Glu Thr Lys  Ser Trp Tyr
    1025                1030                1035

Phe Thr  Glu Asn Met Glu Arg  Asn Cys Arg Gly Ala  Pro Gly Thr
    1040                1045                1050

Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro
    1055                1060                1065

Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1070                1075                1080

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1085                1090                1095

Pro Gly  Ser Ser Pro Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ala
    1100                1105                1110

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1115                1120                1125

Thr Ser  Ser Thr Gly Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala
    1130                1135                1140

Ser Ser  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    1145                1150                1155

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1160                1165                1170

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1175                1180                1185
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Thr Ser  Ser Thr Gly Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala
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Ser Ser  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
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Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser
    1220                 1225                 1230

Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Thr  Pro Gly Ser
    1235                 1240                 1245

Gly Thr  Ala Ser Ser Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1250                 1255                 1260

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1265                 1270                 1275

Pro Gly  Ser Ser Pro Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ser
    1280                 1285                 1290

Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro Gly Ala  Ser Pro Gly
    1295                 1300                 1305

Thr Ser  Ser Thr Gly Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala
    1310                 1315                 1320

Ser Ser  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1325                 1330                 1335

Pro Ala  Ser Ser Ala Pro Cys  Asn Ile Gln Met Glu  Asp Pro Thr
    1340                 1345                 1350

Phe Lys  Glu Asn Tyr Arg Phe  His Ala Ile Asn Gly  Tyr Ile Met
    1355                 1360                 1365

Asp Thr  Leu Pro Gly Leu Val  Met Ala Gln Asp Gln  Arg Ile Arg
    1370                 1375                 1380

Trp Tyr  Leu Leu Ser Met Gly  Ser Asn Glu Asn Ile  His Ser Ile
    1385                 1390                 1395

His Phe  Ser Gly His Val Phe  Thr Val Arg Lys Lys  Glu Glu Tyr
    1400                 1405                 1410

Lys Met  Ala Leu Tyr Asn Leu  Tyr Pro Gly Val Phe  Glu Thr Val
    1415                 1420                 1425

Glu Met  Leu Pro Ser Lys Ala  Gly Ile Trp Arg Val  Glu Cys Leu
    1430                 1435                 1440
```

```
Ile Gly  Glu His Leu His Ala  Gly Met Ser Thr Leu  Phe Leu Val
    1445                 1450                 1455

Tyr Ser  Asn Lys Cys Gln Thr  Pro Leu Gly Met Ala  Ser Gly His
    1460                 1465                 1470

Ile Arg  Asp Phe Gln Ile Thr  Ala Ser Gly Gln Tyr  Gly Gln Trp
    1475                 1480                 1485

Ala Pro  Lys Leu Ala Arg Leu  His Tyr Ser Gly Ser  Ile Asn Ala
    1490                 1495                 1500

Trp Ser  Thr Lys Glu Pro Phe  Ser Trp Ile Lys Val  Asp Leu Leu
    1505                 1510                 1515

Ala Pro  Met Ile Ile His Gly  Ile Lys Thr Gln Gly  Ala Arg Gln
    1520                 1525                 1530

Lys Phe  Ser Ser Leu Tyr Ile  Ser Gln Phe Ile Ile  Met Tyr Ser
    1535                 1540                 1545

Leu Asp  Gly Lys Lys Trp Gln  Thr Tyr Arg Gly Asn  Ser Thr Gly
    1550                 1555                 1560

Thr Leu  Met Val Phe Phe Gly  Asn Val Asp Ser Ser  Gly Ile Lys
    1565                 1570                 1575

His Asn  Ile Phe Asn Pro Pro  Ile Ile Ala Arg Tyr  Ile Arg Leu
    1580                 1585                 1590

His Pro  Thr His Tyr Ser Ile  Arg Ser Thr Leu Arg  Met Glu Leu
    1595                 1600                 1605

Met Gly  Cys Asp Leu Asn Ser  Cys Ser Met Pro Leu  Gly Met Glu
    1610                 1615                 1620

Ser Lys  Ala Ile Ser Asp Ala  Gln Ile Thr Ala Ser  Ser Tyr Phe
    1625                 1630                 1635

Thr Asn  Met Phe Ala Thr Trp  Ser Pro Ser Lys Ala  Arg Leu His
    1640                 1645                 1650

Leu Gln  Gly Arg Ser Asn Ala  Trp Arg Pro Gln Val  Asn Asn Pro
    1655                 1660                 1665

Lys Glu  Trp Leu Gln Val Asp  Phe Gln Lys Thr Met  Lys Val Thr
    1670                 1675                 1680

Gly Val  Thr Thr Gln Gly Val  Lys Ser Leu Leu Thr  Ser Met Tyr
    1685                 1690                 1695
```

```
Val Lys  Glu Phe Leu Ile Ser  Ser Ser Gln Asp Gly  His Gln Trp
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Thr Leu  Phe Phe Gln Asn Gly  Lys Val Lys Val Phe  Gln Gly Asn
    1715                1720                1725

Gln Asp  Ser Phe Thr Pro Val  Val Asn Ser Leu Asp  Pro Pro Leu
    1730                1735                1740

Leu Thr  Arg Tyr Leu Arg Ile  His Pro Gln Ser Trp  Val His Gln
    1745                1750                1755

Ile Ala  Leu Arg Met Glu Val  Leu Gly Cys Glu Ala  Gln Asp Leu
    1760                1765                1770

Tyr Gly  Ala Pro Pro Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr
    1775                1780                1785

Gly Pro  Gly Ser Ser Pro Ser  Ala Ser Thr Gly Thr  Gly Pro Gly
    1790                1795                1800

Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser Pro Gly  Ser Ser Thr
    1805                1810                1815

Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser
    1820                1825                1830

Thr Gly  Thr Gly Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1835                1840                1845

Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly
    1850                1855                1860

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Thr Pro Gly
    1865                1870                1875

Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1880                1885                1890

Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1895                1900                1905

Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala Ser Ser  Ser Ala Ser
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Ser Glu  Gln Lys Leu Ile Ser  Glu Glu Asp Leu Ser  Pro Ala Thr
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Gly His  His His His His His
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atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser

210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
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Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys

485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
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Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Thr Pro Gly
740 745 750

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala

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        755                 760                 765

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    770                 775                 780

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
785                 790                 795                 800

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
                805                 810                 815

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            820                 825                 830

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
        835                 840                 845

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
    850                 855                 860

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
865                 870                 875                 880

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
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1025 1030 1035

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1040 1045 1050

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1055 1060 1065

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1070 1075 1080

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1085 1090 1095

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1100 1105 1110

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1115 1120 1125

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1130 1135 1140

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Gly Ala
1145 1150 1155

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr
1160 1165 1170

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1175 1180 1185

Thr Ser Gly Ser Glu Thr Pro Ala Ser Ser Ala Pro Cys Asn Ile
1190 1195 1200

Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg Phe His Ala
1205 1210 1215

Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu Val Met Ala
1220 1225 1230

Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met Gly Ser Asn
1235 1240 1245

Glu Asn Ile His Ser Ile His Phe Ser Gly His Val Phe Thr Val
1250 1255 1260

Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn Leu Tyr Pro
1265 1270 1275

Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys Ala Gly Ile

1280 1285 1290

Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His Ala Gly Met
1295 1300 1305

Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln Thr Pro Leu
1310 1315 1320

Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile Thr Ala Ser
1325 1330 1335

Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg Leu His Tyr
1340 1345 1350

Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro Phe Ser Trp
1355 1360 1365

Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His Gly Ile Lys
1370 1375 1380

Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr Ile Ser Gln
1385 1390 1395

Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp Gln Thr Tyr
1400 1405 1410

Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe Gly Asn Val
1415 1420 1425

Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro Pro Ile Ile
1430 1435 1440

Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser Ile Arg Ser
1445 1450 1455

Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn Ser Cys Ser
1460 1465 1470

Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp Ala Gln Ile
1475 1480 1485

Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp Ser Pro
1490 1495 1500

Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg
1505 1510 1515

Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln
1520 1525 1530

Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser

1535 1540 1545

Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser
1550 1555 1560

Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val
1565 1570 1575

Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn
1580 1585 1590

Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro
1595 1600 1605

Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly
1610 1615 1620

Cys Glu Ala Gln Asp Leu Tyr Gly Ala Pro Pro Gly Ser Ser Pro
1625 1630 1635

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
1640 1645 1650

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
1655 1660 1665

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
1670 1675 1680

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
1685 1690 1695

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
1700 1705 1710

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
1715 1720 1725

Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
1730 1735 1740

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
1745 1750 1755

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
1760 1765 1770

Ala Ser Ser Ser Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu
1775 1780 1785

Asp Leu Ser Pro Ala Thr Gly His His His His His His

1790 1795 1800

<210> 1446
<211> 5406
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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ctcggtgagc tgcctgtgga cgcaagattt cctcctagag tgccaaaatc ttttccattc 120

aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

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caaagaggaa accagataat gtcagacaag aggaatgtca tcctgttttc tgtatttgat 1740

gagaaccgaa gctggtacct cacagagaat atacaacgct ttctccccaa tccagctgga 1800

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gtttttgata gtttgcagtt gtcagtttgt ttgcatgagg tggcatactg gtacattcta 1920

agcattggag cacagactga cttccttttct gtcttcttct ctggatatac cttcaaacac 1980

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ggcagtgtcc ctcagttcaa gaaagttgtt ttccaggaat ttactgatgg ctcctttact 3000

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gcagaagttg aagataatat catggtaact ttcagaaatc aggcctctcg tccctattcc 3120

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cctgctcatg ggagacaagt gacagtacag gaatttgctc tgtttttcac catctttgat 3420

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cgcatggagt tgatgggctg tgatttaaat agttgcagca tgccattggg aatggagagt 4440

aaagcaatat cagatgcaca gattactgct tcatcctact ttaccaatat gtttgccacc 4500

tggtctcctt caaaagctcg acttcacctc caagggagga gtaatgcctg gagacctcag 4560

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tcgagcacag gcagccccgg aacccctgga tcgggaaccg cgtcgtcaag cgcctcgagc 5340

gaacaaaaac tcatctcaga agaggatctg tcacctgcaa ccggtcatca tcaccatcac 5400

cattga 5406

<210> 1447
<211> 1909
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1447
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

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Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540
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Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Thr Pro Gly
            740                 745                 750

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
        755                 760                 765

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    770                 775                 780

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
785                 790                 795                 800

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
                805                 810                 815
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Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            820                 825                 830

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
        835                 840                 845

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
    850                 855                 860

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
865                 870                 875                 880

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080
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Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                 1090                 1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                 1105                 1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                 1120                 1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                 1135                 1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1145                 1150                 1155

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala
    1160                 1165                 1170

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1175                 1180                 1185

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1190                 1195                 1200

Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
    1205                 1210                 1215

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala
    1220                 1225                 1230

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1235                 1240                 1245

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1250                 1255                 1260

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1265                 1270                 1275

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ala
    1280                 1285                 1290

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Ala Ser  Ser Ala Pro
    1295                 1300                 1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                 1315                 1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                 1330                 1335
```

```
Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1355                1360                1365

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1580                1585                1590
```

```
Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1595                1600                1605

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1610                1615                1620

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1625                1630                1635

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1640                1645                1650

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                1660                1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1670                1675                1680

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1685                1690                1695

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                1705                1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1715                1720                1725

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Pro Gly
    1730                1735                1740

Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly Pro Gly  Ser Ser Pro
    1745                1750                1755

Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Thr Pro Gly  Ser Gly Thr
    1760                1765                1770

Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly
    1775                1780                1785

Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr Gly Thr  Gly Pro Gly
    1790                1795                1800

Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser Pro Gly  Thr Pro Gly
    1805                1810                1815

Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser Ser Thr  Pro Ser Gly
    1820                1825                1830

Ala Thr  Gly Ser Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
    1835                1840                1845
```

Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
1850 1855 1860

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
1865 1870 1875

Ser Gly Thr Ala Ser Ser Ser Ala Ser Ser Glu Gln Lys Leu Ile
1880 1885 1890

Ser Glu Glu Asp Leu Ser Pro Ala Thr Gly His His His His His
1895 1900 1905

His

<210> 1448
<211> 5730
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

agagatctag cttcaggact cattggccct ctcctcatct gctacaaaga atctgtagat 1680

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cctgctcatg ggagacaagt gacagtacag gaatttgctc tgtttttcac catctttgat 3420

gagaccaaaa gctggtactt cactgaaaat atggaaagaa actgcagggg cgcgccaggt 3480

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gcaaccggtc atcatcacca tcaccattga 5730

<210> 1449
<211> 2053
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1449
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser

145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr

420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala

690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Thr Pro Gly
740 745 750

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
755 760 765

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
770 775 780

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
785 790 795 800

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
805 810 815

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
820 825 830

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
835 840 845

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
850 855 860

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
865 870 875 880

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Pro
885 890 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
900 905 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
915 920 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
930 935 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945 950 955 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn

965 970 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
980 985 990

Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu
995 1000 1005

Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val
1010 1015 1020

Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro
1025 1030 1035

Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg
1040 1045 1050

Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
1055 1060 1065

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
1070 1075 1080

Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp
1085 1090 1095

Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val
1100 1105 1110

Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr
1115 1120 1125

Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys
1130 1135 1140

Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Gly Ala
1145 1150 1155

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser
1160 1165 1170

Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
1175 1180 1185

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
1190 1195 1200

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
1205 1210 1215

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala

1220 1225 1230

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser
1235 1240 1245

Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
1250 1255 1260

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
1265 1270 1275

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
1280 1285 1290

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser
1295 1300 1305

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
1310 1315 1320

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
1325 1330 1335

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr
1340 1345 1350

Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
1355 1360 1365

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
1370 1375 1380

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
1385 1390 1395

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala
1400 1405 1410

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser
1415 1420 1425

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
1430 1435 1440

Thr Gly Ser Pro Ala Ser Ser Ala Pro Cys Asn Ile Gln Met Glu
1445 1450 1455

Asp Pro Thr Phe Lys Glu Asn Tyr Arg Phe His Ala Ile Asn Gly
1460 1465 1470

Tyr Ile Met Asp Thr Leu Pro Gly Leu Val Met Ala Gln Asp Gln

1475 1480 1485

Arg Ile Arg Trp Tyr Leu Leu Ser Met Gly Ser Asn Glu Asn Ile
1490 1495 1500

His Ser Ile His Phe Ser Gly His Val Phe Thr Val Arg Lys Lys
1505 1510 1515

Glu Glu Tyr Lys Met Ala Leu Tyr Asn Leu Tyr Pro Gly Val Phe
1520 1525 1530

Glu Thr Val Glu Met Leu Pro Ser Lys Ala Gly Ile Trp Arg Val
1535 1540 1545

Glu Cys Leu Ile Gly Glu His Leu His Ala Gly Met Ser Thr Leu
1550 1555 1560

Phe Leu Val Tyr Ser Asn Lys Cys Gln Thr Pro Leu Gly Met Ala
1565 1570 1575

Ser Gly His Ile Arg Asp Phe Gln Ile Thr Ala Ser Gly Gln Tyr
1580 1585 1590

Gly Gln Trp Ala Pro Lys Leu Ala Arg Leu His Tyr Ser Gly Ser
1595 1600 1605

Ile Asn Ala Trp Ser Thr Lys Glu Pro Phe Ser Trp Ile Lys Val
1610 1615 1620

Asp Leu Leu Ala Pro Met Ile Ile His Gly Ile Lys Thr Gln Gly
1625 1630 1635

Ala Arg Gln Lys Phe Ser Ser Leu Tyr Ile Ser Gln Phe Ile Ile
1640 1645 1650

Met Tyr Ser Leu Asp Gly Lys Lys Trp Gln Thr Tyr Arg Gly Asn
1655 1660 1665

Ser Thr Gly Thr Leu Met Val Phe Phe Gly Asn Val Asp Ser Ser
1670 1675 1680

Gly Ile Lys His Asn Ile Phe Asn Pro Pro Ile Ile Ala Arg Tyr
1685 1690 1695

Ile Arg Leu His Pro Thr His Tyr Ser Ile Arg Ser Thr Leu Arg
1700 1705 1710

Met Glu Leu Met Gly Cys Asp Leu Asn Ser Cys Ser Met Pro Leu
1715 1720 1725

Gly Met Glu Ser Lys Ala Ile Ser Asp Ala Gln Ile Thr Ala Ser

1730 1735 1740

Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp Ser Pro Ser Lys Ala
1745 1750 1755

Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg Pro Gln Val
1760 1765 1770

Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln Lys Thr Met
1775 1780 1785

Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser Leu Leu Thr
1790 1795 1800

Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly
1805 1810 1815

His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val Lys Val Phe
1820 1825 1830

Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn Ser Leu Asp
1835 1840 1845

Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro Gln Ser Trp
1850 1855 1860

Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly Cys Glu Ala
1865 1870 1875

Gln Asp Leu Tyr Gly Ala Pro Pro Gly Ser Ser Pro Ser Ala Ser
1880 1885 1890

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr
1895 1900 1905

Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
1910 1915 1920

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
1925 1930 1935

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser
1940 1945 1950

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
1955 1960 1965

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
1970 1975 1980

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro

1985 1990 1995

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
2000 2005 2010

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
2015 2020 2025

Ser Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
2030 2035 2040

Pro Ala Thr Gly His His His His His His
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<211> 6162
<212> DNA
<213> Artificial Sequence

<220>
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atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

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<210> 1451
<211> 1945
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1451
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

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Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380
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Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655
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Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Pro Gly Ala Ser
            740                 745                 750

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
        755                 760                 765

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
    770                 775                 780

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro
785                 790                 795                 800

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
                805                 810                 815

Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
            820                 825                 830

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
        835                 840                 845

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
    850                 855                 860

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
865                 870                 875                 880

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
                885                 890                 895

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
            900                 905                 910

Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
        915                 920                 925
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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
    930                 935                 940

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
945                 950                 955                 960

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
                965                 970                 975

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
            980                 985                 990

Pro Gly Thr Ser Ser Thr Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser
        995                 1000                1005

Thr Gly  Thr Gly Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
    1010                1015                1020

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Ala Ser
    1025                1030                1035

Ser Pro  Pro Val Leu Lys Arg  His Gln Ala Glu Ile  Thr Arg Thr
    1040                1045                1050

Thr Leu  Gln Ser Asp Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile
    1055                1060                1065

Ser Val  Glu Met Lys Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp
    1070                1075                1080

Glu Asn  Gln Ser Pro Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr
    1085                1090                1095

Phe Ile  Ala Ala Val Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser
    1100                1105                1110

Ser Pro  His Val Leu Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro
    1115                1120                1125

Gln Phe  Lys Lys Val Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe
    1130                1135                1140

Thr Gln  Pro Leu Tyr Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu
    1145                1150                1155

Leu Gly  Pro Tyr Ile Arg Ala  Glu Val Glu Asp Asn  Ile Met Val
    1160                1165                1170

Thr Phe  Arg Asn Gln Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser
    1175                1180                1185
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Leu Ile  Ser Tyr Glu Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg
    1190                 1195                 1200

Lys Asn  Phe Val Lys Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys
    1205                 1210                 1215

Val Gln  His His Met Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys
    1220                 1225                 1230

Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu Glu Lys  Asp Val His
    1235                 1240                 1245

Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys His Thr  Asn Thr Leu
    1250                 1255                 1260

Asn Pro  Ala His Gly Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu
    1265                 1270                 1275

Phe Phe  Thr Ile Phe Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu
    1280                 1285                 1290

Asn Met  Glu Arg Asn Cys Arg  Gly Ala Pro Gly Ser  Pro Ala Gly
    1295                 1300                 1305

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1310                 1315                 1320

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1325                 1330                 1335

Pro Ala  Ser Ser Ala Pro Cys  Asn Ile Gln Met Glu  Asp Pro Thr
    1340                 1345                 1350

Phe Lys  Glu Asn Tyr Arg Phe  His Ala Ile Asn Gly  Tyr Ile Met
    1355                 1360                 1365

Asp Thr  Leu Pro Gly Leu Val  Met Ala Gln Asp Gln  Arg Ile Arg
    1370                 1375                 1380

Trp Tyr  Leu Leu Ser Met Gly  Ser Asn Glu Asn Ile  His Ser Ile
    1385                 1390                 1395

His Phe  Ser Gly His Val Phe  Thr Val Arg Lys Lys  Glu Glu Tyr
    1400                 1405                 1410

Lys Met  Ala Leu Tyr Asn Leu  Tyr Pro Gly Val Phe  Glu Thr Val
    1415                 1420                 1425

Glu Met  Leu Pro Ser Lys Ala  Gly Ile Trp Arg Val  Glu Cys Leu
    1430                 1435                 1440
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Ile Gly  Glu His Leu His Ala  Gly Met Ser Thr Leu  Phe Leu Val
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Tyr Ser  Asn Lys Cys Gln Thr  Pro Leu Gly Met Ala  Ser Gly His
    1460                 1465                 1470

Ile Arg  Asp Phe Gln Ile Thr  Ala Ser Gly Gln Tyr  Gly Gln Trp
    1475                 1480                 1485

Ala Pro  Lys Leu Ala Arg Leu  His Tyr Ser Gly Ser  Ile Asn Ala
    1490                 1495                 1500

Trp Ser  Thr Lys Glu Pro Phe  Ser Trp Ile Lys Val  Asp Leu Leu
    1505                 1510                 1515

Ala Pro  Met Ile Ile His Gly  Ile Lys Thr Gln Gly  Ala Arg Gln
    1520                 1525                 1530

Lys Phe  Ser Ser Leu Tyr Ile  Ser Gln Phe Ile Ile  Met Tyr Ser
    1535                 1540                 1545

Leu Asp  Gly Lys Lys Trp Gln  Thr Tyr Arg Gly Asn  Ser Thr Gly
    1550                 1555                 1560

Thr Leu  Met Val Phe Phe Gly  Asn Val Asp Ser Ser  Gly Ile Lys
    1565                 1570                 1575

His Asn  Ile Phe Asn Pro Pro  Ile Ile Ala Arg Tyr  Ile Arg Leu
    1580                 1585                 1590

His Pro  Thr His Tyr Ser Ile  Arg Ser Thr Leu Arg  Met Glu Leu
    1595                 1600                 1605

Met Gly  Cys Asp Leu Asn Ser  Cys Ser Met Pro Leu  Gly Met Glu
    1610                 1615                 1620

Ser Lys  Ala Ile Ser Asp Ala  Gln Ile Thr Ala Ser  Ser Tyr Phe
    1625                 1630                 1635

Thr Asn  Met Phe Ala Thr Trp  Ser Pro Ser Lys Ala  Arg Leu His
    1640                 1645                 1650

Leu Gln  Gly Arg Ser Asn Ala  Trp Arg Pro Gln Val  Asn Asn Pro
    1655                 1660                 1665

Lys Glu  Trp Leu Gln Val Asp  Phe Gln Lys Thr Met  Lys Val Thr
    1670                 1675                 1680

Gly Val  Thr Thr Gln Gly Val  Lys Ser Leu Leu Thr  Ser Met Tyr
    1685                 1690                 1695
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Val Lys  Glu Phe Leu Ile Ser  Ser Ser Gln Asp Gly  His Gln Trp
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Thr Leu  Phe Phe Gln Asn Gly  Lys Val Lys Val Phe  Gln Gly Asn
    1715                1720                1725

Gln Asp  Ser Phe Thr Pro Val  Val Asn Ser Leu Asp  Pro Pro Leu
    1730                1735                1740

Leu Thr  Arg Tyr Leu Arg Ile  His Pro Gln Ser Trp  Val His Gln
    1745                1750                1755

Ile Ala  Leu Arg Met Glu Val  Leu Gly Cys Glu Ala  Gln Asp Leu
    1760                1765                1770

Tyr Gly  Ala Pro Pro Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr
    1775                1780                1785

Gly Pro  Gly Ser Ser Pro Ser  Ala Ser Thr Gly Thr  Gly Pro Gly
    1790                1795                1800

Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser Pro Gly  Ser Ser Thr
    1805                1810                1815

Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser
    1820                1825                1830

Thr Gly  Thr Gly Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1835                1840                1845

Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly
    1850                1855                1860

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Thr Pro Gly
    1865                1870                1875

Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1880                1885                1890

Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1895                1900                1905

Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala Ser Ser  Ser Ala Ser
    1910                1915                1920

Ser Glu  Gln Lys Leu Ile Ser  Glu Glu Asp Leu Ser  Pro Ala Thr
    1925                1930                1935

Gly His  His His His His His
    1940                1945
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<210> 1452
<211> 5838
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<213> Artificial Sequence

<220>
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atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

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gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

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gtttttgata gtttgcagtt gtcagtttgt ttgcatgagg tggcatactg gtacattcta 1920

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ccatatataa gagcagaagt tgaagataat atcatggtaa ctttcagaaa tcaggcctct 3540

cgtccctatt ccttctattc tagccttatt tcttatgagg aagatcagag gcaaggagca 3600

gaacctagaa aaaactttgt caagcctaat gaaaccaaaa cttacttttg gaaagtgcaa 3660

catcatatgg cacccactaa agatgagttt gactgcaaag cctgggctta tttctctgat 3720

gttgacctgg aaaaagatgt gcactcaggc ctgattggac cccttctggt ctgccacact 3780

aacacactga accctgctca tgggagacaa gtgacagtac aggaatttgc tctgtttttc 3840

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tcacccggca cgtcgagcac aggcagcccc ggaacccctg gatcgggaac cgcgtcgtca 5760

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<211> 2053
<212> PRT
<213> Artificial Sequence

<220>
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20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser

210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys

485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Pro Gly Ala Ser
740 745 750

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser

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        755                 760                 765

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
    770                 775                 780

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro
785                 790                 795                 800

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
                805                 810                 815

Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
            820                 825                 830

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
        835                 840                 845

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
    850                 855                 860

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
865                 870                 875                 880

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
                885                 890                 895

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
            900                 905                 910

Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
        915                 920                 925

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
    930                 935                 940

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
945                 950                 955                 960

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
                965                 970                 975

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
            980                 985                 990

Pro Gly Thr Ser Ser Thr Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser
        995                 1000                1005

Thr Gly  Thr Gly Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
    1010                1015                1020

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Ala Ser
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    1025                1030                1035

Ser Pro  Pro Val Leu Lys Arg  His Gln Ala Glu Ile  Thr Arg Thr
    1040                1045                1050

Thr Leu  Gln Ser Asp Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile
    1055                1060                1065

Ser Val  Glu Met Lys Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp
    1070                1075                1080

Glu Asn  Gln Ser Pro Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr
    1085                1090                1095

Phe Ile  Ala Ala Val Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser
    1100                1105                1110

Ser Pro  His Val Leu Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro
    1115                1120                1125

Gln Phe  Lys Lys Val Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe
    1130                1135                1140

Thr Gln  Pro Leu Tyr Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu
    1145                1150                1155

Leu Gly  Pro Tyr Ile Arg Ala  Glu Val Glu Asp Asn  Ile Met Val
    1160                1165                1170

Thr Phe  Arg Asn Gln Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser
    1175                1180                1185

Leu Ile  Ser Tyr Glu Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg
    1190                1195                1200

Lys Asn  Phe Val Lys Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys
    1205                1210                1215

Val Gln  His His Met Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys
    1220                1225                1230

Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu Glu Lys  Asp Val His
    1235                1240                1245

Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys His Thr  Asn Thr Leu
    1250                1255                1260

Asn Pro  Ala His Gly Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu
    1265                1270                1275

Phe Phe  Thr Ile Phe Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu
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1280 1285 1290

Asn Met Glu Arg Asn Cys Arg Gly Ala Pro Gly Thr Pro Gly Ser
1295 1300 1305

Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
1310 1315 1320

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
1325 1330 1335

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser
1340 1345 1350

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser
1355 1360 1365

Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
1370 1375 1380

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
1385 1390 1395

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser
1400 1405 1410

Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
1415 1420 1425

Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
1430 1435 1440

Thr Gly Ser Pro Ala Ser Ser Ala Pro Cys Asn Ile Gln Met Glu
1445 1450 1455

Asp Pro Thr Phe Lys Glu Asn Tyr Arg Phe His Ala Ile Asn Gly
1460 1465 1470

Tyr Ile Met Asp Thr Leu Pro Gly Leu Val Met Ala Gln Asp Gln
1475 1480 1485

Arg Ile Arg Trp Tyr Leu Leu Ser Met Gly Ser Asn Glu Asn Ile
1490 1495 1500

His Ser Ile His Phe Ser Gly His Val Phe Thr Val Arg Lys Lys
1505 1510 1515

Glu Glu Tyr Lys Met Ala Leu Tyr Asn Leu Tyr Pro Gly Val Phe
1520 1525 1530

Glu Thr Val Glu Met Leu Pro Ser Lys Ala Gly Ile Trp Arg Val

1535 1540 1545

Glu Cys Leu Ile Gly Glu His Leu His Ala Gly Met Ser Thr Leu
1550 1555 1560

Phe Leu Val Tyr Ser Asn Lys Cys Gln Thr Pro Leu Gly Met Ala
1565 1570 1575

Ser Gly His Ile Arg Asp Phe Gln Ile Thr Ala Ser Gly Gln Tyr
1580 1585 1590

Gly Gln Trp Ala Pro Lys Leu Ala Arg Leu His Tyr Ser Gly Ser
1595 1600 1605

Ile Asn Ala Trp Ser Thr Lys Glu Pro Phe Ser Trp Ile Lys Val
1610 1615 1620

Asp Leu Leu Ala Pro Met Ile Ile His Gly Ile Lys Thr Gln Gly
1625 1630 1635

Ala Arg Gln Lys Phe Ser Ser Leu Tyr Ile Ser Gln Phe Ile Ile
1640 1645 1650

Met Tyr Ser Leu Asp Gly Lys Lys Trp Gln Thr Tyr Arg Gly Asn
1655 1660 1665

Ser Thr Gly Thr Leu Met Val Phe Phe Gly Asn Val Asp Ser Ser
1670 1675 1680

Gly Ile Lys His Asn Ile Phe Asn Pro Pro Ile Ile Ala Arg Tyr
1685 1690 1695

Ile Arg Leu His Pro Thr His Tyr Ser Ile Arg Ser Thr Leu Arg
1700 1705 1710

Met Glu Leu Met Gly Cys Asp Leu Asn Ser Cys Ser Met Pro Leu
1715 1720 1725

Gly Met Glu Ser Lys Ala Ile Ser Asp Ala Gln Ile Thr Ala Ser
1730 1735 1740

Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp Ser Pro Ser Lys Ala
1745 1750 1755

Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg Pro Gln Val
1760 1765 1770

Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln Lys Thr Met
1775 1780 1785

Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser Leu Leu Thr

1790 1795 1800

Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly
1805 1810 1815

His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val Lys Val Phe
1820 1825 1830

Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn Ser Leu Asp
1835 1840 1845

Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro Gln Ser Trp
1850 1855 1860

Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly Cys Glu Ala
1865 1870 1875

Gln Asp Leu Tyr Gly Ala Pro Pro Gly Ser Ser Pro Ser Ala Ser
1880 1885 1890

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr
1895 1900 1905

Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
1910 1915 1920

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
1925 1930 1935

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser
1940 1945 1950

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
1955 1960 1965

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
1970 1975 1980

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
1985 1990 1995

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
2000 2005 2010

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
2015 2020 2025

Ser Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser
2030 2035 2040

Pro Ala Thr Gly His His His His His His

2045 2050

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<211> 6162
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

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ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

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Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445
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Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720
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```
Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Pro Gly Ala Ser
            740                 745                 750

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
        755                 760                 765

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
    770                 775                 780

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro
785                 790                 795                 800

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
                805                 810                 815

Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
            820                 825                 830

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
        835                 840                 845

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
    850                 855                 860

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
865                 870                 875                 880

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
                885                 890                 895

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
            900                 905                 910

Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
        915                 920                 925

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
    930                 935                 940

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
945                 950                 955                 960

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
                965                 970                 975

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
            980                 985                 990
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Pro Gly Thr Ser Ser Thr Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser
       995                 1000                1005

Thr Gly  Thr Gly Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
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Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Ala Ser
    1025                1030                1035

Ser Pro  Pro Val Leu Lys Arg  His Gln Ala Glu Ile  Thr Arg Thr
    1040                1045                1050

Thr Leu  Gln Ser Asp Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile
    1055                1060                1065

Ser Val  Glu Met Lys Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp
    1070                1075                1080

Glu Asn  Gln Ser Pro Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr
    1085                1090                1095

Phe Ile  Ala Ala Val Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser
    1100                1105                1110

Ser Pro  His Val Leu Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro
    1115                1120                1125

Gln Phe  Lys Lys Val Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe
    1130                1135                1140

Thr Gln  Pro Leu Tyr Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu
    1145                1150                1155

Leu Gly  Pro Tyr Ile Arg Ala  Glu Val Glu Asp Asn  Ile Met Val
    1160                1165                1170

Thr Phe  Arg Asn Gln Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser
    1175                1180                1185

Leu Ile  Ser Tyr Glu Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg
    1190                1195                1200

Lys Asn  Phe Val Lys Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys
    1205                1210                1215

Val Gln  His His Met Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys
    1220                1225                1230

Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu Glu Lys  Asp Val His
    1235                1240                1245
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Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys His Thr  Asn Thr Leu
    1250                1255                1260

Asn Pro  Ala His Gly Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu
    1265                1270                1275

Phe Phe  Thr Ile Phe Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu
    1280                1285                1290

Asn Met  Glu Arg Asn Cys Arg  Gly Ala Pro Gly Thr  Pro Gly Ser
    1295                1300                1305

Gly Thr  Ala Ser Ser Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1310                1315                1320

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1325                1330                1335

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser
    1340                1345                1350

Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro Gly Ala  Ser Pro Gly
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Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1370                1375                1380

Thr Gly  Ser Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1385                1390                1395

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1400                1405                1410

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1415                1420                1425

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
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Thr Gly  Ser Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1445                1450                1455

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Thr
    1460                1465                1470

Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro
    1475                1480                1485

Ser Gly  Ala Thr Gly Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala
    1490                1495                1500
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Ser Ser  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
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Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser
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Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro Gly Ser  Ser Pro Ser
    1535                 1540                 1545

Ala Ser  Thr Gly Thr Gly Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1550                 1555                 1560

Thr Gly  Ser Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1565                 1570                 1575

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Ala Ser
    1580                 1585                 1590

Ser Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1595                 1600                 1605

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1610                 1615                 1620

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1625                 1630                 1635

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1640                 1645                 1650

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1655                 1660                 1665

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1670                 1675                 1680

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1685                 1690                 1695

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1700                 1705                 1710

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1715                 1720                 1725

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1730                 1735                 1740

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1745                 1750                 1755
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Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1760                1765                1770

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1775                1780                1785

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1790                1795                1800

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1805                1810                1815

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1820                1825                1830

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1835                1840                1845

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1850                1855                1860

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1865                1870                1875

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1880                1885                1890

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1895                1900                1905

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1910                1915                1920

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1925                1930                1935

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1940                1945                1950

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1955                1960                1965

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1970                1975                1980

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    1985                1990                1995

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    2000                2005                2010
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Pro Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
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Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
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Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
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Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr
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Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
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Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
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Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
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Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
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Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Ala Ser Ser Glu Gln
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His His His His
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu

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            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
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450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala

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Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Ser Pro Ala
            740                 745                 750

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
        755                 760                 765

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
    770                 775                 780

Ala Ser Ser Pro Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg
785                 790                 795                 800

Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
                805                 810                 815

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu
            820                 825                 830

Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile
        835                 840                 845

Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His
    850                 855                 860

Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys
865                 870                 875                 880

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
                885                 890                 895

Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
            900                 905                 910

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser
        915                 920                 925

Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln
    930                 935                 940

Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
945                 950                 955                 960

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp
                965                 970                 975

Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu
            980                 985                 990

Lys Asp Val His Ser Gly Leu Ile  Gly Pro Leu Leu Val  Cys His Thr
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Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr Val Gln Glu
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Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys Ser Trp Tyr
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Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Gly Ala Pro Gly Ser
1040 1045 1050

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
1055 1060 1065

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
1070 1075 1080

Ser Glu Thr Pro Ala Ser Ser Ala Pro Cys Asn Ile Gln Met Glu
1085 1090 1095

Asp Pro Thr Phe Lys Glu Asn Tyr Arg Phe His Ala Ile Asn Gly
1100 1105 1110

Tyr Ile Met Asp Thr Leu Pro Gly Leu Val Met Ala Gln Asp Gln
1115 1120 1125

Arg Ile Arg Trp Tyr Leu Leu Ser Met Gly Ser Asn Glu Asn Ile
1130 1135 1140

His Ser Ile His Phe Ser Gly His Val Phe Thr Val Arg Lys Lys
1145 1150 1155

Glu Glu Tyr Lys Met Ala Leu Tyr Asn Leu Tyr Pro Gly Val Phe
1160 1165 1170

Glu Thr Val Glu Met Leu Pro Ser Lys Ala Gly Ile Trp Arg Val
1175 1180 1185

Glu Cys Leu Ile Gly Glu His Leu His Ala Gly Met Ser Thr Leu
1190 1195 1200

Phe Leu Val Tyr Ser Asn Lys Cys Gln Thr Pro Leu Gly Met Ala
1205 1210 1215

Ser Gly His Ile Arg Asp Phe Gln Ile Thr Ala Ser Gly Gln Tyr
1220 1225 1230

Gly Gln Trp Ala Pro Lys Leu Ala Arg Leu His Tyr Ser Gly Ser
1235 1240 1245

Ile Asn Ala Trp Ser Thr Lys Glu Pro Phe Ser Trp Ile Lys Val

1250 1255 1260

Asp Leu Leu Ala Pro Met Ile Ile His Gly Ile Lys Thr Gln Gly
1265 1270 1275

Ala Arg Gln Lys Phe Ser Ser Leu Tyr Ile Ser Gln Phe Ile Ile
1280 1285 1290

Met Tyr Ser Leu Asp Gly Lys Lys Trp Gln Thr Tyr Arg Gly Asn
1295 1300 1305

Ser Thr Gly Thr Leu Met Val Phe Phe Gly Asn Val Asp Ser Ser
1310 1315 1320

Gly Ile Lys His Asn Ile Phe Asn Pro Pro Ile Ile Ala Arg Tyr
1325 1330 1335

Ile Arg Leu His Pro Thr His Tyr Ser Ile Arg Ser Thr Leu Arg
1340 1345 1350

Met Glu Leu Met Gly Cys Asp Leu Asn Ser Cys Ser Met Pro Leu
1355 1360 1365

Gly Met Glu Ser Lys Ala Ile Ser Asp Ala Gln Ile Thr Ala Ser
1370 1375 1380

Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp Ser Pro Ser Lys Ala
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Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg Pro Gln Val
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Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln Lys Thr Met
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Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser Leu Leu Thr
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Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly
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His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val Lys Val Phe
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Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn Ser Leu Asp
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Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro Gln Ser Trp
1490 1495 1500

Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly Cys Glu Ala

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Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
1790 1795 1800

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Ala Ser Ser Glu Gln
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Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala Thr Gly His His
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His His His His
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

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Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445
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Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720
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Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Ser Pro Ala
            740                 745                 750

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
        755                 760                 765

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
    770                 775                 780

Ala Ser Ser Pro Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg
785                 790                 795                 800

Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
                805                 810                 815

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu
            820                 825                 830

Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile
        835                 840                 845

Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His
    850                 855                 860

Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys
865                 870                 875                 880

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
                885                 890                 895

Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
            900                 905                 910

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser
        915                 920                 925

Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln
    930                 935                 940

Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
945                 950                 955                 960

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp
                965                 970                 975

Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu
            980                 985                 990
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Lys Asp Val His Ser Gly Leu Ile  Gly Pro Leu Leu Val  Cys His Thr
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Asn Thr  Leu Asn Pro Ala His  Gly Arg Gln Val Thr  Val Gln Glu
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Phe Ala  Leu Phe Phe Thr Ile  Phe Asp Glu Thr Lys  Ser Trp Tyr
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Phe Thr  Glu Asn Met Glu Arg  Asn Cys Arg Gly Ala  Pro Gly Thr
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Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro Gly Ala  Ser Pro Gly
    1055                1060                1065


Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1070                1075                1080


Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    1085                1090                1095


Pro Gly  Ser Ser Pro Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Thr
    1100                1105                1110


Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro Gly Ala  Ser Pro Gly
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Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
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Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
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Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser
    1160                1165                1170


Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1175                1180                1185


Thr Ser  Ser Thr Gly Ser Pro  Ala Ser Ser Ala Pro  Cys Asn Ile
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Gln Met  Glu Asp Pro Thr Phe  Lys Glu Asn Tyr Arg  Phe His Ala
    1205                1210                1215


Ile Asn  Gly Tyr Ile Met Asp  Thr Leu Pro Gly Leu  Val Met Ala
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Gln Asp  Gln Arg Ile Arg Trp  Tyr Leu Leu Ser Met  Gly Ser Asn
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Glu Asn  Ile His Ser Ile His  Phe Ser Gly His Val  Phe Thr Val
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Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
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Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    1280                 1285                 1290

Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    1295                 1300                 1305

Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    1310                 1315                 1320

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
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Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
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Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
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Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
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Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
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Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
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Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    1415                 1420                 1425

Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
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Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1445                 1450                 1455

Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
    1460                 1465                 1470

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
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Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
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Ser Lys  Ala Arg Leu His Leu  Gln Gly Arg Ser Asn  Ala Trp Arg
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Pro Gln  Val Asn Asn Pro Lys  Glu Trp Leu Gln Val  Asp Phe Gln
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Lys Thr  Met Lys Val Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser
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Leu Leu  Thr Ser Met Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser
    1550                 1555                 1560

Gln Asp  Gly His Gln Trp Thr  Leu Phe Phe Gln Asn  Gly Lys Val
    1565                 1570                 1575

Lys Val  Phe Gln Gly Asn Gln  Asp Ser Phe Thr Pro  Val Val Asn
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Ser Leu  Asp Pro Pro Leu Leu  Thr Arg Tyr Leu Arg  Ile His Pro
    1595                 1600                 1605

Gln Ser  Trp Val His Gln Ile  Ala Leu Arg Met Glu  Val Leu Gly
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Cys Glu  Ala Gln Asp Leu Tyr  Gly Ala Pro Pro Gly  Ala Ser Pro
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Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
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Ser Thr  Gly Ser Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
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Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly
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Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser Pro Gly  Ser Ser Thr
    1685                 1690                 1695

Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Thr Pro Gly  Ser Gly Thr
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Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly
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Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly
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Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly Pro Gly  Ser Ser Pro
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Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser
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1865 1870 1875

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
1880 1885 1890

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
1895 1900 1905

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Ala Ser
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285
```

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Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560
```

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Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Ser Pro Ala
            740                 745                 750

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
        755                 760                 765

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
    770                 775                 780

Ala Ser Ser Pro Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg
785                 790                 795                 800

Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
                805                 810                 815

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu
            820                 825                 830
```

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Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile
        835                 840                 845

Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His
    850                 855                 860

Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys
865                 870                 875                 880

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
                885                 890                 895

Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
            900                 905                 910

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser
        915                 920                 925

Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln
    930                 935                 940

Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
945                 950                 955                 960

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp
                965                 970                 975

Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu
            980                 985                 990

Lys Asp Val His Ser Gly Leu Ile  Gly Pro Leu Leu Val  Cys His Thr
        995                 1000                1005

Asn Thr  Leu Asn Pro Ala His  Gly Arg Gln Val Thr  Val Gln Glu
    1010                1015                1020

Phe Ala  Leu Phe Phe Thr Ile  Phe Asp Glu Thr Lys  Ser Trp Tyr
    1025                1030                1035

Phe Thr  Glu Asn Met Glu Arg  Asn Cys Arg Gly Ala  Pro Gly Thr
    1040                1045                1050

Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro
    1055                1060                1065

Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1070                1075                1080

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1085                1090                1095
```

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Pro Gly  Ser Ser Pro Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ala
    1100                 1105                 1110

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1115                 1120                 1125

Thr Ser  Ser Thr Gly Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala
    1130                 1135                 1140

Ser Ser  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    1145                 1150                 1155

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1160                 1165                 1170

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1175                 1180                 1185

Thr Ser  Ser Thr Gly Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala
    1190                 1195                 1200

Ser Ser  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1205                 1210                 1215

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser
    1220                 1225                 1230

Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Thr  Pro Gly Ser
    1235                 1240                 1245

Gly Thr  Ala Ser Ser Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1250                 1255                 1260

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1265                 1270                 1275

Pro Gly  Ser Ser Pro Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ser
    1280                 1285                 1290

Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro Gly Ala  Ser Pro Gly
    1295                 1300                 1305

Thr Ser  Ser Thr Gly Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala
    1310                 1315                 1320

Ser Ser  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1325                 1330                 1335

Pro Ala  Ser Ser Ala Pro Cys  Asn Ile Gln Met Glu  Asp Pro Thr
    1340                 1345                 1350
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```
Phe Lys  Glu Asn Tyr Arg Phe  His Ala Ile Asn Gly  Tyr Ile Met
    1355                1360                1365

Asp Thr  Leu Pro Gly Leu Val  Met Ala Gln Asp Gln  Arg Ile Arg
    1370                1375                1380

Trp Tyr  Leu Leu Ser Met Gly  Ser Asn Glu Asn Ile  His Ser Ile
    1385                1390                1395

His Phe  Ser Gly His Val Phe  Thr Val Arg Lys Lys  Glu Glu Tyr
    1400                1405                1410

Lys Met  Ala Leu Tyr Asn Leu  Tyr Pro Gly Val Phe  Glu Thr Val
    1415                1420                1425

Glu Met  Leu Pro Ser Lys Ala  Gly Ile Trp Arg Val  Glu Cys Leu
    1430                1435                1440

Ile Gly  Glu His Leu His Ala  Gly Met Ser Thr Leu  Phe Leu Val
    1445                1450                1455

Tyr Ser  Asn Lys Cys Gln Thr  Pro Leu Gly Met Ala  Ser Gly His
    1460                1465                1470

Ile Arg  Asp Phe Gln Ile Thr  Ala Ser Gly Gln Tyr  Gly Gln Trp
    1475                1480                1485

Ala Pro  Lys Leu Ala Arg Leu  His Tyr Ser Gly Ser  Ile Asn Ala
    1490                1495                1500

Trp Ser  Thr Lys Glu Pro Phe  Ser Trp Ile Lys Val  Asp Leu Leu
    1505                1510                1515

Ala Pro  Met Ile Ile His Gly  Ile Lys Thr Gln Gly  Ala Arg Gln
    1520                1525                1530

Lys Phe  Ser Ser Leu Tyr Ile  Ser Gln Phe Ile Ile  Met Tyr Ser
    1535                1540                1545

Leu Asp  Gly Lys Lys Trp Gln  Thr Tyr Arg Gly Asn  Ser Thr Gly
    1550                1555                1560

Thr Leu  Met Val Phe Phe Gly  Asn Val Asp Ser Ser  Gly Ile Lys
    1565                1570                1575

His Asn  Ile Phe Asn Pro Pro  Ile Ile Ala Arg Tyr  Ile Arg Leu
    1580                1585                1590

His Pro  Thr His Tyr Ser Ile  Arg Ser Thr Leu Arg  Met Glu Leu
    1595                1600                1605
```

```
Met Gly  Cys Asp Leu Asn Ser  Cys Ser Met Pro Leu  Gly Met Glu
    1610                 1615                 1620

Ser Lys  Ala Ile Ser Asp Ala  Gln Ile Thr Ala Ser  Ser Tyr Phe
    1625                 1630                 1635

Thr Asn  Met Phe Ala Thr Trp  Ser Pro Ser Lys Ala  Arg Leu His
    1640                 1645                 1650

Leu Gln  Gly Arg Ser Asn Ala  Trp Arg Pro Gln Val  Asn Asn Pro
    1655                 1660                 1665

Lys Glu  Trp Leu Gln Val Asp  Phe Gln Lys Thr Met  Lys Val Thr
    1670                 1675                 1680

Gly Val  Thr Thr Gln Gly Val  Lys Ser Leu Leu Thr  Ser Met Tyr
    1685                 1690                 1695

Val Lys  Glu Phe Leu Ile Ser  Ser Ser Gln Asp Gly  His Gln Trp
    1700                 1705                 1710

Thr Leu  Phe Phe Gln Asn Gly  Lys Val Lys Val Phe  Gln Gly Asn
    1715                 1720                 1725

Gln Asp  Ser Phe Thr Pro Val  Val Asn Ser Leu Asp  Pro Pro Leu
    1730                 1735                 1740

Leu Thr  Arg Tyr Leu Arg Ile  His Pro Gln Ser Trp  Val His Gln
    1745                 1750                 1755

Ile Ala  Leu Arg Met Glu Val  Leu Gly Cys Glu Ala  Gln Asp Leu
    1760                 1765                 1770

Tyr Gly  Ala Pro Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1775                 1780                 1785

Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly
    1790                 1795                 1800

Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser Pro Gly  Ser Ser Thr
    1805                 1810                 1815

Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Thr Pro Gly  Ser Gly Thr
    1820                 1825                 1830

Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly
    1835                 1840                 1845

Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly
    1850                 1855                 1860
```

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Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr
    1865                 1870                 1875

Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser
    1880                 1885                 1890

Thr Gly  Thr Gly Pro Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr
    1895                 1900                 1905

Gly Pro  Gly Ala Ser Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly
    1910                 1915                 1920

Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser Pro Gly  Ser Ser Thr
    1925                 1930                 1935

Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser
    1940                 1945                 1950

Thr Gly  Thr Gly Pro Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr
    1955                 1960                 1965

Gly Pro  Gly Ala Ser Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly
    1970                 1975                 1980

Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser Pro Gly  Ser Ser Thr
    1985                 1990                 1995

Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser
    2000                 2005                 2010

Thr Gly  Thr Gly Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    2015                 2020                 2025

Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr Gly Thr  Gly Pro Gly
    2030                 2035                 2040

Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser Pro Gly  Ser Ser Thr
    2045                 2050                 2055

Pro Ser  Gly Ala Thr Gly Ser  Ala Ser Ser Glu Gln  Lys Leu Ile
    2060                 2065                 2070

Ser Glu  Glu Asp Leu Ser Pro  Ala Thr Gly His His  His His His
    2075                 2080                 2085

His
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<212> PRT
<213> Artificial Sequence

<220>
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

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Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445
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Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720
```

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Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Thr Pro Gly
            740                 745                 750

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
        755                 760                 765

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    770                 775                 780

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
785                 790                 795                 800

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
                805                 810                 815

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            820                 825                 830

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
        835                 840                 845

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
    850                 855                 860

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
865                 870                 875                 880

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990
```

```
Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
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Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
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Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1145                1150                1155

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr
    1160                1165                1170

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1175                1180                1185

Thr Ser  Gly Ser Glu Thr Pro  Ala Ser Ser Ala Pro  Cys Asn Ile
    1190                1195                1200

Gln Met  Glu Asp Pro Thr Phe  Lys Glu Asn Tyr Arg  Phe His Ala
    1205                1210                1215

Ile Asn  Gly Tyr Ile Met Asp  Thr Leu Pro Gly Leu  Val Met Ala
    1220                1225                1230

Gln Asp  Gln Arg Ile Arg Trp  Tyr Leu Leu Ser Met  Gly Ser Asn
    1235                1240                1245
```

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Glu Asn  Ile His Ser Ile His  Phe Ser Gly His Val  Phe Thr Val
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Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
    1265                 1270                 1275

Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    1280                 1285                 1290

Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    1295                 1300                 1305

Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    1310                 1315                 1320

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
    1325                 1330                 1335

Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
    1340                 1345                 1350

Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
    1355                 1360                 1365

Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
    1370                 1375                 1380

Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    1385                 1390                 1395

Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
    1400                 1405                 1410

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    1415                 1420                 1425

Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
    1430                 1435                 1440

Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    1445                 1450                 1455

Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
    1460                 1465                 1470

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    1475                 1480                 1485

Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
    1490                 1495                 1500
```

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Ser Lys  Ala Arg Leu His Leu  Gln Gly Arg Ser Asn  Ala Trp Arg
    1505                 1510                 1515

Pro Gln  Val Asn Asn Pro Lys  Glu Trp Leu Gln Val  Asp Phe Gln
    1520                 1525                 1530

Lys Thr  Met Lys Val Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser
    1535                 1540                 1545

Leu Leu  Thr Ser Met Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser
    1550                 1555                 1560

Gln Asp  Gly His Gln Trp Thr  Leu Phe Phe Gln Asn  Gly Lys Val
    1565                 1570                 1575

Lys Val  Phe Gln Gly Asn Gln  Asp Ser Phe Thr Pro  Val Val Asn
    1580                 1585                 1590

Ser Leu  Asp Pro Pro Leu Leu  Thr Arg Tyr Leu Arg  Ile His Pro
    1595                 1600                 1605

Gln Ser  Trp Val His Gln Ile  Ala Leu Arg Met Glu  Val Leu Gly
    1610                 1615                 1620

Cys Glu  Ala Gln Asp Leu Tyr  Gly Ala Pro Pro Gly  Ala Ser Pro
    1625                 1630                 1635

Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
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Ser Thr  Gly Ser Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
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Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly
    1670                 1675                 1680

Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser Pro Gly  Ser Ser Thr
    1685                 1690                 1695

Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Thr Pro Gly  Ser Gly Thr
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Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly
    1715                 1720                 1725

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly
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Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly Pro Gly  Ser Ser Pro
    1745                 1750                 1755
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Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ala Ser Pro  Gly Thr Ser
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Ser Thr  Gly Ser Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
    1775                1780                1785

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly
    1790                1795                1800

Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly Pro Gly  Ser Ser Pro
    1805                1810                1815

Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ala Ser Pro  Gly Thr Ser
    1820                1825                1830

Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
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Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly
    1850                1855                1860

Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly Pro Gly  Ala Ser Pro
    1865                1870                1875

Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser
    1880                1885                1890

Thr Gly  Thr Gly Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
    1895                1900                1905

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Ala Ser
    1910                1915                1920

Ser Glu  Gln Lys Leu Ile Ser  Glu Glu Asp Leu Ser  Pro Ala Thr
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Gly His  His His His His His
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<400> 1464

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285
```

```
Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560
```

```
Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Thr Pro Gly
            740                 745                 750

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
        755                 760                 765

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    770                 775                 780

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
785                 790                 795                 800

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
                805                 810                 815

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            820                 825                 830
```

```
Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
        835                 840                 845

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
    850                 855                 860

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
865                 870                 875                 880

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095
```

```
Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
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Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1145                1150                1155

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala
    1160                1165                1170

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1175                1180                1185

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1190                1195                1200

Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
    1205                1210                1215

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala
    1220                1225                1230

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1235                1240                1245

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1250                1255                1260

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1265                1270                1275

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ala
    1280                1285                1290

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Ala Ser  Ser Ala Pro
    1295                1300                1305

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1310                1315                1320

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1325                1330                1335

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1340                1345                1350
```

```
Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
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Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1370                1375                1380

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1385                1390                1395

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1400                1405                1410

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1415                1420                1425

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1430                1435                1440

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1445                1450                1455

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1460                1465                1470

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1475                1480                1485

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1490                1495                1500

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1505                1510                1515

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1520                1525                1530

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1535                1540                1545

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1550                1555                1560

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1565                1570                1575

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
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Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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```

```
Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
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Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
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Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
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Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1655                 1660                 1665

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
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Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
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Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1700                 1705                 1710

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
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Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Pro Gly
    1730                 1735                 1740

Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser Pro Gly  Ala Ser Pro
    1745                 1750                 1755

Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Thr Pro Gly  Ser Gly Thr
    1760                 1765                 1770

Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly
    1775                 1780                 1785

Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly
    1790                 1795                 1800

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Thr Pro Gly
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Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser Ser Thr  Pro Ser Gly
    1820                 1825                 1830

Ala Thr  Gly Ser Pro Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly
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Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr Gly Thr  Gly Pro Gly
    1850                 1855                 1860
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Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
1865 1870 1875

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
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Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
1895 1900 1905

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
1910 1915 1920

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
1925 1930 1935

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
1940 1945 1950

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
1955 1960 1965

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
1970 1975 1980

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
1985 1990 1995

Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr
2000 2005 2010

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240
```

```
Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510
```

```
Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Gly Thr Pro Gly
            740                 745                 750

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
        755                 760                 765

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    770                 775                 780
```

```
Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
785                 790                 795                 800

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
                805                 810                 815

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            820                 825                 830

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
        835                 840                 845

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
    850                 855                 860

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
865                 870                 875                 880

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
        915                 920                 925

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
    930                 935                 940

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu
945                 950                 955                 960

Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn
                965                 970                 975

Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
            980                 985                 990

Glu Phe Thr Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
        995                 1000                1005

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1010                1015                1020

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1025                1030                1035

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1040                1045                1050
```

```
Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1055                1060                1065

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1070                1075                1080

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1085                1090                1095

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1100                1105                1110

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1115                1120                1125

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1130                1135                1140

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1145                1150                1155

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser
    1160                1165                1170

Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Ser  Ser Thr Pro
    1175                1180                1185

Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1190                1195                1200

Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
    1205                1210                1215

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1220                1225                1230

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Thr  Pro Gly Ser
    1235                1240                1245

Gly Thr  Ala Ser Ser Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1250                1255                1260

Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    1265                1270                1275

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1280                1285                1290

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Thr  Pro Gly Ser
    1295                1300                1305
```

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Gly Thr  Ala Ser Ser Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
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Thr Gly  Ser Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
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Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Thr
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Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro
    1355                 1360                 1365

Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1370                 1375                 1380

Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
    1385                 1390                 1395

Pro Gly  Ser Ser Pro Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ala
    1400                 1405                 1410

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Thr  Pro Gly Ser
    1415                 1420                 1425

Gly Thr  Ala Ser Ser Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1430                 1435                 1440

Thr Gly  Ser Pro Ala Ser Ser  Ala Pro Cys Asn Ile  Gln Met Glu
    1445                 1450                 1455

Asp Pro  Thr Phe Lys Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly
    1460                 1465                 1470

Tyr Ile  Met Asp Thr Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln
    1475                 1480                 1485

Arg Ile  Arg Trp Tyr Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile
    1490                 1495                 1500

His Ser  Ile His Phe Ser Gly  His Val Phe Thr Val  Arg Lys Lys
    1505                 1510                 1515

Glu Glu  Tyr Lys Met Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe
    1520                 1525                 1530

Glu Thr  Val Glu Met Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val
    1535                 1540                 1545

Glu Cys  Leu Ile Gly Glu His  Leu His Ala Gly Met  Ser Thr Leu
    1550                 1555                 1560
```

```
Phe Leu  Val Tyr Ser Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala
    1565                1570                1575

Ser Gly  His Ile Arg Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr
    1580                1585                1590

Gly Gln  Trp Ala Pro Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser
    1595                1600                1605

Ile Asn  Ala Trp Ser Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val
    1610                1615                1620

Asp Leu  Leu Ala Pro Met Ile  Ile His Gly Ile Lys  Thr Gln Gly
    1625                1630                1635

Ala Arg  Gln Lys Phe Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile
    1640                1645                1650

Met Tyr  Ser Leu Asp Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn
    1655                1660                1665

Ser Thr  Gly Thr Leu Met Val  Phe Phe Gly Asn Val  Asp Ser Ser
    1670                1675                1680

Gly Ile  Lys His Asn Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr
    1685                1690                1695

Ile Arg  Leu His Pro Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg
    1700                1705                1710

Met Glu  Leu Met Gly Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu
    1715                1720                1725

Gly Met  Glu Ser Lys Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser
    1730                1735                1740

Ser Tyr  Phe Thr Asn Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala
    1745                1750                1755

Arg Leu  His Leu Gln Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val
    1760                1765                1770

Asn Asn  Pro Lys Glu Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met
    1775                1780                1785

Lys Val  Thr Gly Val Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr
    1790                1795                1800

Ser Met  Tyr Val Lys Glu Phe  Leu Ile Ser Ser Ser  Gln Asp Gly
    1805                1810                1815
```

```
His Gln  Trp Thr Leu Phe Phe  Gln Asn Gly Lys Val  Lys Val Phe
    1820                1825                1830

Gln Gly  Asn Gln Asp Ser Phe  Thr Pro Val Val Asn  Ser Leu Asp
    1835                1840                1845

Pro Pro  Leu Leu Thr Arg Tyr  Leu Arg Ile His Pro  Gln Ser Trp
    1850                1855                1860

Val His  Gln Ile Ala Leu Arg  Met Glu Val Leu Gly  Cys Glu Ala
    1865                1870                1875

Gln Asp  Leu Tyr Gly Ala Pro  Pro Gly Ala Ser Pro  Gly Thr Ser
    1880                1885                1890

Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1895                1900                1905

Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly
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Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Thr Pro Gly
    1925                1930                1935

Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser Ser Thr  Pro Ser Gly
    1940                1945                1950

Ala Thr  Gly Ser Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
    1955                1960                1965

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly
    1970                1975                1980

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Pro
    1985                1990                1995

Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ser Ser Pro  Ser Ala Ser
    2000                2005                2010

Thr Gly  Thr Gly Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    2015                2020                2025

Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly
    2030                2035                2040

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Pro
    2045                2050                2055

Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ser Ser Pro  Ser Ala Ser
    2060                2065                2070
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Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
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Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
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Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
2105 2110 2115

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser
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Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr
2135 2140 2145

Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
2150 2155 2160

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Ala Ser Ser Glu Gln
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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255
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Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525
```

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Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Pro Gly Ala Ser
            740                 745                 750

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
        755                 760                 765

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
    770                 775                 780

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro
785                 790                 795                 800
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Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
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Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
            820                 825                 830

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
        835                 840                 845

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
    850                 855                 860

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
865                 870                 875                 880

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
                885                 890                 895

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
            900                 905                 910

Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
        915                 920                 925

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
    930                 935                 940

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
945                 950                 955                 960

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
                965                 970                 975

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
            980                 985                 990

Pro Gly Thr Ser Ser Thr Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser
        995                 1000                1005

Thr Gly  Thr Gly Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
    1010                1015                1020

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Ala Ser
    1025                1030                1035

Ser Pro  Pro Val Leu Lys Arg  His Gln Ala Glu Ile  Thr Arg Thr
    1040                1045                1050

Thr Leu  Gln Ser Asp Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile
    1055                1060                1065
```

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Ser Val  Glu Met Lys Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp
    1070                1075                1080

Glu Asn  Gln Ser Pro Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr
    1085                1090                1095

Phe Ile  Ala Ala Val Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser
    1100                1105                1110

Ser Pro  His Val Leu Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro
    1115                1120                1125

Gln Phe  Lys Lys Val Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe
    1130                1135                1140

Thr Gln  Pro Leu Tyr Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu
    1145                1150                1155

Leu Gly  Pro Tyr Ile Arg Ala  Glu Val Glu Asp Asn  Ile Met Val
    1160                1165                1170

Thr Phe  Arg Asn Gln Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser
    1175                1180                1185

Leu Ile  Ser Tyr Glu Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg
    1190                1195                1200

Lys Asn  Phe Val Lys Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys
    1205                1210                1215

Val Gln  His His Met Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys
    1220                1225                1230

Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu Glu Lys  Asp Val His
    1235                1240                1245

Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys His Thr  Asn Thr Leu
    1250                1255                1260

Asn Pro  Ala His Gly Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu
    1265                1270                1275

Phe Phe  Thr Ile Phe Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu
    1280                1285                1290

Asn Met  Glu Arg Asn Cys Arg  Gly Ala Pro Gly Ser  Pro Ala Gly
    1295                1300                1305

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1310                1315                1320
```

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Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
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Pro Ala  Ser Ser Ala Pro Cys  Asn Ile Gln Met Glu  Asp Pro Thr
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Phe Lys  Glu Asn Tyr Arg Phe  His Ala Ile Asn Gly  Tyr Ile Met
    1355                 1360                 1365

Asp Thr  Leu Pro Gly Leu Val  Met Ala Gln Asp Gln  Arg Ile Arg
    1370                 1375                 1380

Trp Tyr  Leu Leu Ser Met Gly  Ser Asn Glu Asn Ile  His Ser Ile
    1385                 1390                 1395

His Phe  Ser Gly His Val Phe  Thr Val Arg Lys Lys  Glu Glu Tyr
    1400                 1405                 1410

Lys Met  Ala Leu Tyr Asn Leu  Tyr Pro Gly Val Phe  Glu Thr Val
    1415                 1420                 1425

Glu Met  Leu Pro Ser Lys Ala  Gly Ile Trp Arg Val  Glu Cys Leu
    1430                 1435                 1440

Ile Gly  Glu His Leu His Ala  Gly Met Ser Thr Leu  Phe Leu Val
    1445                 1450                 1455

Tyr Ser  Asn Lys Cys Gln Thr  Pro Leu Gly Met Ala  Ser Gly His
    1460                 1465                 1470

Ile Arg  Asp Phe Gln Ile Thr  Ala Ser Gly Gln Tyr  Gly Gln Trp
    1475                 1480                 1485

Ala Pro  Lys Leu Ala Arg Leu  His Tyr Ser Gly Ser  Ile Asn Ala
    1490                 1495                 1500

Trp Ser  Thr Lys Glu Pro Phe  Ser Trp Ile Lys Val  Asp Leu Leu
    1505                 1510                 1515

Ala Pro  Met Ile Ile His Gly  Ile Lys Thr Gln Gly  Ala Arg Gln
    1520                 1525                 1530

Lys Phe  Ser Ser Leu Tyr Ile  Ser Gln Phe Ile Ile  Met Tyr Ser
    1535                 1540                 1545

Leu Asp  Gly Lys Lys Trp Gln  Thr Tyr Arg Gly Asn  Ser Thr Gly
    1550                 1555                 1560

Thr Leu  Met Val Phe Phe Gly  Asn Val Asp Ser Ser  Gly Ile Lys
    1565                 1570                 1575
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His Asn  Ile Phe Asn Pro Pro  Ile Ile Ala Arg Tyr  Ile Arg Leu
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His Pro  Thr His Tyr Ser Ile  Arg Ser Thr Leu Arg  Met Glu Leu
    1595                 1600                 1605

Met Gly  Cys Asp Leu Asn Ser  Cys Ser Met Pro Leu  Gly Met Glu
    1610                 1615                 1620

Ser Lys  Ala Ile Ser Asp Ala  Gln Ile Thr Ala Ser  Ser Tyr Phe
    1625                 1630                 1635

Thr Asn  Met Phe Ala Thr Trp  Ser Pro Ser Lys Ala  Arg Leu His
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Leu Gln  Gly Arg Ser Asn Ala  Trp Arg Pro Gln Val  Asn Asn Pro
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Lys Glu  Trp Leu Gln Val Asp  Phe Gln Lys Thr Met  Lys Val Thr
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Gly Val  Thr Thr Gln Gly Val  Lys Ser Leu Leu Thr  Ser Met Tyr
    1685                 1690                 1695

Val Lys  Glu Phe Leu Ile Ser  Ser Ser Gln Asp Gly  His Gln Trp
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Thr Leu  Phe Phe Gln Asn Gly  Lys Val Lys Val Phe  Gln Gly Asn
    1715                 1720                 1725

Gln Asp  Ser Phe Thr Pro Val  Val Asn Ser Leu Asp  Pro Pro Leu
    1730                 1735                 1740

Leu Thr  Arg Tyr Leu Arg Ile  His Pro Gln Ser Trp  Val His Gln
    1745                 1750                 1755

Ile Ala  Leu Arg Met Glu Val  Leu Gly Cys Glu Ala  Gln Asp Leu
    1760                 1765                 1770

Tyr Gly  Ala Pro Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1775                 1780                 1785

Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly
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Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser Pro Gly  Ser Ser Thr
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Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Thr Pro Gly  Ser Gly Thr
    1820                 1825                 1830
```

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Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly
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Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly
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Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr
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Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser
    1880                 1885                 1890

Thr Gly  Thr Gly Pro Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr
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Gly Pro  Gly Ala Ser Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly
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Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser Pro Gly  Ser Ser Thr
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Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser
    1940                 1945                 1950

Thr Gly  Thr Gly Pro Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr
    1955                 1960                 1965

Gly Pro  Gly Ala Ser Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly
    1970                 1975                 1980

Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser Pro Gly  Ser Ser Thr
    1985                 1990                 1995

Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser
    2000                 2005                 2010

Thr Gly  Thr Gly Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    2015                 2020                 2025

Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr Gly Thr  Gly Pro Gly
    2030                 2035                 2040

Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser Pro Gly  Ser Ser Thr
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Pro Ser  Gly Ala Thr Gly Ser  Ala Ser Ser Glu Gln  Lys Leu Ile
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Ser Glu  Glu Asp Leu Ser Pro  Ala Thr Gly His His  His His His
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His

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

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Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415
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Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685
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Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Pro Gly Ala Ser
            740                 745                 750

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
        755                 760                 765

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
    770                 775                 780

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro
785                 790                 795                 800

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
                805                 810                 815

Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
            820                 825                 830

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
        835                 840                 845

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
    850                 855                 860

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
865                 870                 875                 880

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
                885                 890                 895

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
            900                 905                 910

Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
        915                 920                 925

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
    930                 935                 940

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
945                 950                 955                 960
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Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
                965                 970                 975

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
            980                 985                 990

Pro Gly Thr Ser Ser Thr Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser
        995                 1000                1005

Thr Gly  Thr Gly Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
    1010                1015                1020

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Ala Ser
    1025                1030                1035

Ser Pro  Pro Val Leu Lys Arg  His Gln Ala Glu Ile  Thr Arg Thr
    1040                1045                1050

Thr Leu  Gln Ser Asp Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile
    1055                1060                1065

Ser Val  Glu Met Lys Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp
    1070                1075                1080

Glu Asn  Gln Ser Pro Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr
    1085                1090                1095

Phe Ile  Ala Ala Val Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser
    1100                1105                1110

Ser Pro  His Val Leu Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro
    1115                1120                1125

Gln Phe  Lys Lys Val Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe
    1130                1135                1140

Thr Gln  Pro Leu Tyr Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu
    1145                1150                1155

Leu Gly  Pro Tyr Ile Arg Ala  Glu Val Glu Asp Asn  Ile Met Val
    1160                1165                1170

Thr Phe  Arg Asn Gln Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser
    1175                1180                1185

Leu Ile  Ser Tyr Glu Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg
    1190                1195                1200

Lys Asn  Phe Val Lys Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys
    1205                1210                1215
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Val Gln  His His Met Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys
    1220                 1225                 1230

Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu Glu Lys  Asp Val His
    1235                 1240                 1245

Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys His Thr  Asn Thr Leu
    1250                 1255                 1260

Asn Pro  Ala His Gly Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu
    1265                 1270                 1275

Phe Phe  Thr Ile Phe Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu
    1280                 1285                 1290

Asn Met  Glu Arg Asn Cys Arg  Gly Ala Pro Gly Thr  Pro Gly Ser
    1295                 1300                 1305

Gly Thr  Ala Ser Ser Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1310                 1315                 1320

Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    1325                 1330                 1335

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ser
    1340                 1345                 1350

Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro Gly Thr  Pro Gly Ser
    1355                 1360                 1365

Gly Thr  Ala Ser Ser Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
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Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    1385                 1390                 1395

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ser
    1400                 1405                 1410

Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Ser  Ser Thr Pro
    1415                 1420                 1425

Ser Gly  Ala Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1430                 1435                 1440

Thr Gly  Ser Pro Ala Ser Ser  Ala Pro Cys Asn Ile  Gln Met Glu
    1445                 1450                 1455

Asp Pro  Thr Phe Lys Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly
    1460                 1465                 1470
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Tyr Ile  Met Asp Thr Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln
    1475                 1480                 1485

Arg Ile  Arg Trp Tyr Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile
    1490                 1495                 1500

His Ser  Ile His Phe Ser Gly  His Val Phe Thr Val  Arg Lys Lys
    1505                 1510                 1515

Glu Glu  Tyr Lys Met Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe
    1520                 1525                 1530

Glu Thr  Val Glu Met Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val
    1535                 1540                 1545

Glu Cys  Leu Ile Gly Glu His  Leu His Ala Gly Met  Ser Thr Leu
    1550                 1555                 1560

Phe Leu  Val Tyr Ser Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala
    1565                 1570                 1575

Ser Gly  His Ile Arg Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr
    1580                 1585                 1590

Gly Gln  Trp Ala Pro Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser
    1595                 1600                 1605

Ile Asn  Ala Trp Ser Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val
    1610                 1615                 1620

Asp Leu  Leu Ala Pro Met Ile  Ile His Gly Ile Lys  Thr Gln Gly
    1625                 1630                 1635

Ala Arg  Gln Lys Phe Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile
    1640                 1645                 1650

Met Tyr  Ser Leu Asp Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn
    1655                 1660                 1665

Ser Thr  Gly Thr Leu Met Val  Phe Phe Gly Asn Val  Asp Ser Ser
    1670                 1675                 1680

Gly Ile  Lys His Asn Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr
    1685                 1690                 1695

Ile Arg  Leu His Pro Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg
    1700                 1705                 1710

Met Glu  Leu Met Gly Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu
    1715                 1720                 1725
```

```
Gly Met  Glu Ser Lys Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser
    1730                1735                1740

Ser Tyr  Phe Thr Asn Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala
    1745                1750                1755

Arg Leu  His Leu Gln Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val
    1760                1765                1770

Asn Asn  Pro Lys Glu Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met
    1775                1780                1785

Lys Val  Thr Gly Val Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr
    1790                1795                1800

Ser Met  Tyr Val Lys Glu Phe  Leu Ile Ser Ser Ser  Gln Asp Gly
    1805                1810                1815

His Gln  Trp Thr Leu Phe Phe  Gln Asn Gly Lys Val  Lys Val Phe
    1820                1825                1830

Gln Gly  Asn Gln Asp Ser Phe  Thr Pro Val Val Asn  Ser Leu Asp
    1835                1840                1845

Pro Pro  Leu Leu Thr Arg Tyr  Leu Arg Ile His Pro  Gln Ser Trp
    1850                1855                1860

Val His  Gln Ile Ala Leu Arg  Met Glu Val Leu Gly  Cys Glu Ala
    1865                1870                1875

Gln Asp  Leu Tyr Gly Ala Pro  Pro Gly Ala Ser Pro  Gly Thr Ser
    1880                1885                1890

Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1895                1900                1905

Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly
    1910                1915                1920

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Thr Pro Gly
    1925                1930                1935

Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser Ser Thr  Pro Ser Gly
    1940                1945                1950

Ala Thr  Gly Ser Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
    1955                1960                1965

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly
    1970                1975                1980
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Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
1985 1990 1995

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
2000 2005 2010

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
2015 2020 2025

Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
2030 2035 2040

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
2045 2050 2055

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
2060 2065 2070

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
2075 2080 2085

Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
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Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
2105 2110 2115

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser
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Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr
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Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
2150 2155 2160

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Ala Ser Ser Glu Gln
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Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala Thr Gly His His
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His His His His
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20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

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His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430
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Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700
```

```
Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ala Pro Pro Gly Ala Ser
            740                 745                 750

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
        755                 760                 765

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
    770                 775                 780

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro
785                 790                 795                 800

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
                805                 810                 815

Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
            820                 825                 830

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
        835                 840                 845

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
    850                 855                 860

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
865                 870                 875                 880

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
                885                 890                 895

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
            900                 905                 910

Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
        915                 920                 925

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
    930                 935                 940

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
945                 950                 955                 960

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
                965                 970                 975
```

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Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
            980                 985                 990

Pro Gly Thr Ser Ser Thr Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser
        995                 1000                1005

Thr Gly  Thr Gly Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
    1010                1015                1020

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Ala Ser
    1025                1030                1035

Ser Pro  Pro Val Leu Lys Arg  His Gln Ala Glu Ile  Thr Arg Thr
    1040                1045                1050

Thr Leu  Gln Ser Asp Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile
    1055                1060                1065

Ser Val  Glu Met Lys Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp
    1070                1075                1080

Glu Asn  Gln Ser Pro Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr
    1085                1090                1095

Phe Ile  Ala Ala Val Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser
    1100                1105                1110

Ser Pro  His Val Leu Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro
    1115                1120                1125

Gln Phe  Lys Lys Val Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe
    1130                1135                1140

Thr Gln  Pro Leu Tyr Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu
    1145                1150                1155

Leu Gly  Pro Tyr Ile Arg Ala  Glu Val Glu Asp Asn  Ile Met Val
    1160                1165                1170

Thr Phe  Arg Asn Gln Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser
    1175                1180                1185

Leu Ile  Ser Tyr Glu Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg
    1190                1195                1200

Lys Asn  Phe Val Lys Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys
    1205                1210                1215

Val Gln  His His Met Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys
    1220                1225                1230
```

```
Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu Glu Lys  Asp Val His
    1235                1240                1245

Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys His Thr  Asn Thr Leu
    1250                1255                1260

Asn Pro  Ala His Gly Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu
    1265                1270                1275

Phe Phe  Thr Ile Phe Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu
    1280                1285                1290

Asn Met  Glu Arg Asn Cys Arg  Gly Ala Pro Gly Thr  Pro Gly Ser
    1295                1300                1305

Gly Thr  Ala Ser Ser Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1310                1315                1320

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1325                1330                1335

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser
    1340                1345                1350

Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro Gly Ala  Ser Pro Gly
    1355                1360                1365

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1370                1375                1380

Thr Gly  Ser Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1385                1390                1395

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    1400                1405                1410

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1415                1420                1425

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1430                1435                1440

Thr Gly  Ser Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1445                1450                1455

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Thr
    1460                1465                1470

Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro
    1475                1480                1485
```

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Ser Gly  Ala Thr Gly Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala
    1490                 1495                 1500

Ser Ser  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1505                 1510                 1515

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser
    1520                 1525                 1530

Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro Gly Ser  Ser Pro Ser
    1535                 1540                 1545

Ala Ser  Thr Gly Thr Gly Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1550                 1555                 1560

Thr Gly  Ser Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1565                 1570                 1575

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Ala Ser
    1580                 1585                 1590

Ser Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1595                 1600                 1605

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1610                 1615                 1620

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1625                 1630                 1635

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1640                 1645                 1650

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1655                 1660                 1665

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1670                 1675                 1680

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1685                 1690                 1695

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1700                 1705                 1710

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1715                 1720                 1725

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1730                 1735                 1740
```

```
Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1745                1750                1755

Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1760                1765                1770

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1775                1780                1785

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1790                1795                1800

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1805                1810                1815

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1820                1825                1830

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1835                1840                1845

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1850                1855                1860

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1865                1870                1875

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1880                1885                1890

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1895                1900                1905

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1910                1915                1920

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1925                1930                1935

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1940                1945                1950

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1955                1960                1965

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1970                1975                1980

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    1985                1990                1995
```

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Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    2000                2005                2010

Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Ala
    2015                2020                2025

Pro Pro  Gly Ala Ser Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly
    2030                2035                2040

Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser Pro Gly  Thr Pro Gly
    2045                2050                2055

Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser Ser Thr  Pro Ser Gly
    2060                2065                2070

Ala Thr  Gly Ser Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
    2075                2080                2085

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly
    2090                2095                2100

Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser Pro Gly  Ser Ser Thr
    2105                2110                2115

Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser Ser Thr  Pro Ser Gly
    2120                2125                2130

Ala Thr  Gly Ser Pro Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr
    2135                2140                2145

Gly Pro  Gly Ser Ser Pro Ser  Ala Ser Thr Gly Thr  Gly Pro Gly
    2150                2155                2160

Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser Pro Gly  Thr Pro Gly
    2165                2170                2175

Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser Ser Thr  Pro Ser Gly
    2180                2185                2190

Ala Thr  Gly Ser Pro Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr
    2195                2200                2205

Gly Pro  Gly Ser Ser Pro Ser  Ala Ser Thr Gly Thr  Gly Pro Gly
    2210                2215                2220

Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser Pro Gly  Ala Ser Pro
    2225                2230                2235

Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ser Ser Thr  Pro Ser Gly
    2240                2245                2250
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Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr
2255 2260 2265

Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
2270 2275 2280

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
2285 2290 2295

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
2300 2305 2310

Ala Thr Gly Ser Ala Ser Ser Glu Gln Lys Leu Ile Ser Glu Glu
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Asp Leu Ser Pro Ala Thr Gly His His His His His His
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tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

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catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

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<212> PRT
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<400> 1475

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240
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Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510
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Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
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Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780
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Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
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Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
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Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
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Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
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Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
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Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
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Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
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Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
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Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1085                 1090                 1095

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
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Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
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Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
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Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
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Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
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Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
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Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
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Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
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Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
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Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
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Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1250                 1255                 1260

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
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Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
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Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
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Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
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Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
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Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
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Ser His His His His His His His His
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<210> 1477
<211> 1449
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1477

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            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255
```

```
Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525
```

```
Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800
```

```
Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1025                1030                1035

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1040                1045                1050

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1055                1060                1065
```

```
Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1070                1075                1080

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1085                1090                1095

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1100                1105                1110

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1115                1120                1125

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1130                1135                1140

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1145                1150                1155

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1160                1165                1170

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1175                1180                1185

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1190                1195                1200

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1205                1210                1215

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1220                1225                1230

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1235                1240                1245

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1250                1255                1260

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1265                1270                1275

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1280                1285                1290

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1295                1300                1305

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1310                1315                1320
```

```
Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
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Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1340                 1345                 1350

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1355                 1360                 1365

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1370                 1375                 1380

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1385                 1390                 1395

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1400                 1405                 1410

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1415                 1420                 1425

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Ala Ser  Ser His His
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His His  His His His His
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<210> 1478
<211> 4350
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 1478

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac      180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt      240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct      300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa      360

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<210> 1479
<211> 1899
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 1479

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Gly Ala Pro Gly Ser Ser
            20                  25                  30

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
        35                  40                  45

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
    50                  55                  60

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
65                  70                  75                  80

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
                85                  90                  95

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
            100                 105                 110

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
        115                 120                 125

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
    130                 135                 140

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
145                 150                 155                 160

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Ala Ser Ser
                165                 170                 175

Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr
            180                 185                 190

Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu
        195                 200                 205

Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro
    210                 215                 220

Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn
225                 230                 235                 240

Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp
                245                 250                 255

Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu
            260                 265                 270
```

```
Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp
        275                 280                 285

Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu
    290                 295                 300

Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser
305                 310                 315                 320

Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys
                325                 330                 335

Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe
            340                 345                 350

Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
        355                 360                 365

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val
    370                 375                 380

Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg
385                 390                 395                 400

Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val
                405                 410                 415

His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg
            420                 425                 430

Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr
        435                 440                 445

Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser
    450                 455                 460

His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro
465                 470                 475                 480

Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr
                485                 490                 495

Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp
            500                 505                 510

Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His
        515                 520                 525

Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp
    530                 535                 540
```

```
Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln
545                 550                 555                 560

Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val
                565                 570                 575

Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile
            580                 585                 590

Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
        595                 600                 605

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn
    610                 615                 620

Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
625                 630                 635                 640

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
                645                 650                 655

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
            660                 665                 670

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn
        675                 680                 685

Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys
    690                 695                 700

Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys
705                 710                 715                 720

Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr
                725                 730                 735

Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
            740                 745                 750

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn
        755                 760                 765

Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val
    770                 775                 780

Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser
785                 790                 795                 800

Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp
                805                 810                 815
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Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met
            820                 825                 830

Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
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Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn
    850                 855                 860

Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
865                 870                 875                 880

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro
                885                 890                 895

Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln
            900                 905                 910

Gly Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
        915                 920                 925

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
    930                 935                 940

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
945                 950                 955                 960

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
                965                 970                 975

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly
            980                 985                 990

Thr Ser Ser Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
        995                 1000                1005

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1010                1015                1020

Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser
    1025                1030                1035

Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    1040                1045                1050

Ser Ser  Thr Gly Ser Pro Ala  Ser Ser Ser Asp Gln  Glu Glu Ile
    1055                1060                1065

Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met Lys Lys  Glu Asp Phe
    1070                1075                1080
```

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Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser Pro Arg  Ser Phe Gln
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Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala Val Glu  Arg Leu Trp
    1100                 1105                 1110

Asp Tyr  Gly Met Ser Ser Ser  Pro His Val Leu Arg  Asn Arg Ala
    1115                 1120                 1125

Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys Val Val  Phe Gln Glu
    1130                 1135                 1140

Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu Tyr Arg  Gly Glu Leu
    1145                 1150                 1155

Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val
    1160                 1165                 1170

Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro
    1175                 1180                 1185

Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg
    1190                 1195                 1200

Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr
    1205                 1210                 1215

Lys Thr  Tyr Phe Trp Lys Val  Gln His His Met Ala  Pro Thr Lys
    1220                 1225                 1230

Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp
    1235                 1240                 1245

Leu Glu  Lys Asp Val His Ser  Gly Leu Ile Gly Pro  Leu Leu Val
    1250                 1255                 1260

Cys His  Thr Asn Thr Leu Asn  Pro Ala His Gly Arg  Gln Val Thr
    1265                 1270                 1275

Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys
    1280                 1285                 1290

Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Gly Ala
    1295                 1300                 1305

Pro Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    1310                 1315                 1320

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1325                 1330                 1335
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Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1340                1345                1350

Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1355                1360                1365

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1370                1375                1380

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Pro  Ala Gly Ser
    1385                1390                1395

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1400                1405                1410

Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1415                1420                1425

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1430                1435                1440

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ala Ser  Ser Ala Pro
    1445                1450                1455

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1460                1465                1470

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1475                1480                1485

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1490                1495                1500

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1505                1510                1515

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1520                1525                1530

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1535                1540                1545

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1550                1555                1560

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1565                1570                1575

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1580                1585                1590
```

```
Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1595                1600                1605

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1610                1615                1620

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1625                1630                1635

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1640                1645                1650

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1655                1660                1665

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1670                1675                1680

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1685                1690                1695

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1700                1705                1710

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1715                1720                1725

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1730                1735                1740

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1745                1750                1755

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1760                1765                1770

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1775                1780                1785

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1790                1795                1800

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1805                1810                1815

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1820                1825                1830

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1835                1840                1845
```

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1850 1855 1860

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1865 1870 1875

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Ala Ser Ser His His
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His His His His His His
1895

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ccaggttcgt caccctcagc atcaacgggg acgggaccgg gggcgtcacc aggaacgtcc 180

tccaccggct cgccgggtgc atcacccgga acgtcatcga ccggatcgcc agggagctcg 240

acgccatcag gcgcaacagg atcacctggc tcaagcccta gcgcgtcaac cggcacgggt 300

ccgggtgcct cccctggcac gtccagcacc ggatcacccg gatcgagccc atccgcctca 360

accggaaccg gacccggtac accagggtcg ggaacagcct cctcgtcacc aggctcctca 420

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<211> 1596
<212> PRT
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<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1481

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu

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            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
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    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
```

725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
740 745 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
755 760 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
770 775 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785 790 795 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
805 810 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
820 825 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
835 840 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
850 855 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865 870 875 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
885 890 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
900 905 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
915 920 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
930 935 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945 950 955 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
965 970 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
980 985 990

Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro

995 1000 1005

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1010 1015 1020

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1025 1030 1035

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1040 1045 1050

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1055 1060 1065

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1070 1075 1080

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1085 1090 1095

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1100 1105 1110

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1115 1120 1125

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1130 1135 1140

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1145 1150 1155

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1160 1165 1170

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1175 1180 1185

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1190 1195 1200

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1205 1210 1215

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1220 1225 1230

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1235 1240 1245

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser

```
    1250                1255                1260

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1265                1270                1275

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1280                1285                1290

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1295                1300                1305

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1310                1315                1320

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1325                1330                1335

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1340                1345                1350

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1355                1360                1365

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1370                1375                1380

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1385                1390                1395

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1400                1405                1410

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1415                1420                1425

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Thr Ser
    1430                1435                1440

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala
    1445                1450                1455

Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1460                1465                1470

Ser Gly  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1475                1480                1485

Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu
    1490                1495                1500

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser
```

1505 1510 1515

Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
1520 1525 1530

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
1535 1540 1545

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser
1550 1555 1560

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro
1565 1570 1575

Ser Glu Gly Ser Ala Pro Gly Ala Ser Ser His His His His His
1580 1585 1590

His His His
1595

<210> 1482
<211> 4791
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 1482
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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960

gaagcttatg tcaaagtaga cagctgtcca gaggaacccc aactacgaat gaaaaataat 1020

gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080

gatgatgaca actctccttc ctttatccaa attcgctcag ttgccaagaa gcatcctaaa 1140

acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

gcccccgatg acagaagtta taaaagtcaa tatttgaaca atggccctca gcggattggt 1260

aggaagtaca aaaaagtccg atttatggca tacacagatg aaacctttaa gactcgtgaa 1320

gctattcagc atgaatcagg aatcttggga cctttacttt atggggaagt tggagacaca 1380

ctgttgatta tatttaagaa tcaagcaagc agaccatata acatctaccc tcacggaatc 1440

actgatgtcc gtcctttgta ttcaaggaga ttaccaaaag gtgtaaaaca tttgaaggat 1500

tttccaattc tgccaggaga aatattcaaa tataaatgga cagtgactgt agaagatggg 1560

ccaactaaat cagatcctcg gtgcctgacc cgctattact ctagtttcgt taatatggag 1620

agagatctag cttcaggact cattggccct ctcctcatct gctacaaaga atctgtagat 1680

caaagaggaa accagataat gtcagacaag aggaatgtca tcctgttttc tgtatttgat 1740

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gtgcagcttg aggatccaga gttccaagcc tccaacatca tgcacagcat caatggctat 1860

gtttttgata gtttgcagtt gtcagtttgt ttgcatgagg tggcatactg gtacattcta 1920

agcattggag cacagactga cttccttttct gtcttcttct ctggatatac cttcaaacac 1980

aaaatggtct atgaagacac actcacccta ttcccattct caggagaaac tgtcttcatg 2040

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ggcatgaccg ccttactgaa ggtttctagt tgtgacaaga acactggtga ttattacgag 2160

gacagttatg aagatatttc agcatacttg ctgagtaaaa acaatgccat tgaaccaaga 2220

agcttctctc aaaacccacc agtcttgaaa cgccatcaac gggaaataac tcgtactact 2280

cttcagtcag atcaagagga aatcgattat gatgatacca tatcagttga aatgaagaag 2340

gaagattttg acatttatga tgaggatgaa aatcagagcc cccgcagctt tcaaaagaaa 2400

acacgacact attttattgc tgcagtggag aggctctggg attatgggat gagtagctcc 2460

ccacatgttc taagaaacag ggctcagagt ggcagtgtcc ctcagttcaa gaaagttgtt 2520

ttccaggaat ttactgatgg ctcctttact cagcccttat accgtggaga actaaatgaa 2580

catttgggac tcctggggcc atatataaga gcagaagttg aagataatat catggtaact 2640

ttcagaaatc aggcctctcg tccctattcc ttctattcta gccttatttc ttatgaggaa 2700

gatcagaggc aaggagcaga acctagaaaa aactttgtca agcctaatga aaccaaaact 2760

tacttttgga aagtgcaaca tcatatggca cccactaaag atgagtttga ctgcaaagcc 2820

tgggcttatt tctctgatgt tgacctggaa aaagatgtgc actcaggcct gattggaccc 2880

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gaatttgctc tgtttttcac catctttgat gagaccaaaa gctggtactt cactgaaaat 3000

atggaaagaa actgcagggc tccctgcaat atccagatgg aagatcccac ttttaaagag 3060

aattatcgct tccatgcaat caatggctac ataatggata cactacctgg cttagtaatg 3120

gctcaggatc aaaggattcg atggtatctg ctcagcatgg gcagcaatga aaacatccat 3180

tctattcatt tcagtggaca tgtgttcact gtacgaaaaa aagaggagta taaaatggca 3240

ctgtacaatc tctatccagg tgtttttgag acagtggaaa tgttaccatc caaagctgga 3300

atttggcggg tggaatgcct tattggcgag catctacatg ctgggatgag cacacttttt 3360

ctggtgtaca gcaataagtg tcagactccc ctgggaatgg cttctggaca cattagagat 3420

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tattccggat caatcaatgc ctggagcacc aaggagccct tttcttggat caaggtggat 3540

ctgttggcac caatgattat tcacggcatc aagacccagg gtgcccgtca gaagttctcc 3600

agcctctaca tctctcagtt tatcatcatg tatagtcttg atgggaagaa gtggcagact 3660

tatcgaggaa attccactgg aaccttaatg gtcttctttg gcaatgtgga ttcatctggg 3720

ataaaacaca atatttttaa ccctccaatt attgctcgat acatccgttt gcacccaact 3780

cattatagca ttcgcagcac tcttcgcatg gagttgatgg gctgtgattt aaatagttgc 3840

agcatgccat tgggaatgga gagtaaagca atatcagatg cacagattac tgcttcatcc 3900

tactttacca atatgtttgc cacctggtct ccttcaaaag ctcgacttca cctccaaggg 3960

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cagaagacaa tgaaagtcac aggagtaact actcagggag taaaatctct gcttaccagc 4080

atgtatgtga aggagttcct catctccagc agtcaagatg gccatcagtg gactctcttt 4140

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caccagattg ccctgaggat ggaggttctg ggctgcgagg cacaggacct ctacggcgcg 4320

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ggctcagaac ccgctacctc aggctccgag acacccggct ctcctgctgg gagtcccact 4560

tccaccgagg aaggaacatc cactgagcct agtgagggct ctgcccctgg aaccagcaca 4620

gagccaagtg agggcagtgc accaggatcc gagccagcaa ccagcgggtc cgagactccc 4680

gggacctctg agtctgccac cccagagagc ggacccggca cttcaaccga gccctccgaa 4740

ggatcagcac caggtgcctc gagccatcac catcatcacc atcaccattg a 4791

<210> 1483
<211> 1596
<212> PRT
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1483

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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255
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Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525
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Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800
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```
Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1025                1030                1035

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1040                1045                1050

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1055                1060                1065
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Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1070                1075                1080

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1085                1090                1095

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1100                1105                1110

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1115                1120                1125

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1130                1135                1140

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1145                1150                1155

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1160                1165                1170

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1175                1180                1185

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1190                1195                1200

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1205                1210                1215

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1220                1225                1230

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1235                1240                1245

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1250                1255                1260

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1265                1270                1275

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1280                1285                1290

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1295                1300                1305

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1310                1315                1320
```

```
Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1325                1330                1335

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1340                1345                1350

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1355                1360                1365

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1370                1375                1380

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1385                1390                1395

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1400                1405                1410

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1415                1420                1425

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Pro Gly
    1430                1435                1440

Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly Pro Gly  Ser Ser Pro
    1445                1450                1455

Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Thr Pro Gly  Ser Gly Thr
    1460                1465                1470

Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly
    1475                1480                1485

Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr Gly Thr  Gly Pro Gly
    1490                1495                1500

Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser Pro Gly  Thr Pro Gly
    1505                1510                1515

Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser Ser Thr  Pro Ser Gly
    1520                1525                1530

Ala Thr  Gly Ser Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
    1535                1540                1545

Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly
    1550                1555                1560

Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser Pro Gly  Thr Pro Gly
    1565                1570                1575
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Ser Gly Thr Ala Ser Ser Ser Ala Ser Ser His His His His His His
1580 1585 1590

His His His
1595

<210> 1484
<211> 4791
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 1484
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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180

atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240

tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300

gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360

agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420

ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480

catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540

gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact acttttgct 600

gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660

gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720

tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780

ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840

catcgccagg ctagcttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900

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Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

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Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
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Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335
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Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605
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Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
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Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
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Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1025                1030                1035

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1040                1045                1050

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1055                1060                1065

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1070                1075                1080

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1085                1090                1095

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1100                1105                1110

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1115                1120                1125

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
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Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
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Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1160                1165                1170

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1175                1180                1185

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
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Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
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Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1220                1225                1230

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1235                1240                1245

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
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Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
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Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
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Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
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Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
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Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
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Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1355                1360                1365

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
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Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
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Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
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Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1415                 1420                 1425

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Gly Thr
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1445                 1450                 1455

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1460                 1465                 1470

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
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Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1505                 1510                 1515

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1520                 1525                 1530

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1535                 1540                 1545

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
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Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
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Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    1580                 1585                 1590

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
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Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
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Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
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Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
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Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
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Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala
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Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
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<210> 1487
<211> 1740
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1487
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Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

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His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480
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Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750
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```
Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020
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Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
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Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1040                1045                1050

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1055                1060                1065

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1070                1075                1080

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1085                1090                1095

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1100                1105                1110

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1115                1120                1125

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1130                1135                1140

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1145                1150                1155

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1160                1165                1170

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1175                1180                1185

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1190                1195                1200

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1205                1210                1215

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1220                1225                1230

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1235                1240                1245

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1250                1255                1260

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1265                1270                1275
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Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1280                 1285                 1290

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1295                 1300                 1305

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1310                 1315                 1320

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1325                 1330                 1335

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1340                 1345                 1350

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1355                 1360                 1365

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1370                 1375                 1380

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1385                 1390                 1395

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1400                 1405                 1410

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1415                 1420                 1425

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Ala  Pro Pro Gly
    1430                 1435                 1440

Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser Pro Gly  Ala Ser Pro
    1445                 1450                 1455

Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Thr Pro Gly  Ser Gly Thr
    1460                 1465                 1470

Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly
    1475                 1480                 1485

Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly
    1490                 1495                 1500

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Thr Pro Gly
    1505                 1510                 1515

Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser Ser Thr  Pro Ser Gly
    1520                 1525                 1530
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Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
1535 1540 1545

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
1550 1555 1560

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
1565 1570 1575

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
1580 1585 1590

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
1595 1600 1605

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
1610 1615 1620

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
1625 1630 1635

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
1640 1645 1650

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
1655 1660 1665

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
1670 1675 1680

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
1685 1690 1695

Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr
1700 1705 1710

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
1715 1720 1725

Ser Ala Ser Ser His His His His His His His His
1730 1735 1740

<210> 1488
<211> 5223
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polynucleotide

<400> 1488

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aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac      180
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tga 5223

<210> 1489
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1489
Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
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Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro
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<210> 1490
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1490
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
1 5 10 15

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro
35

<210> 1491
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1491
Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
1 5 10 15

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
20 25 30

Thr Gly Ser Pro
35

<210> 1492
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1492
Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
1 5 10 15

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala
20 25 30

Thr Gly Ser Pro
35

<210> 1493
<211> 42
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic

polypeptide

<400> 1493
Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro
1 5 10 15

Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
20 25 30

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
35 40

<210> 1494
<211> 42
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1494
Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro
1 5 10 15

Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
20 25 30

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
35 40

<210> 1495
<211> 42
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1495
Gly Ala Pro Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
1 5 10 15

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
20 25 30

Ala Thr Gly Ser Pro Gly Pro Ser Gly Pro
35 40

<210> 1496
<211> 42
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1496
Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser
1 5 10 15

Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala
20 25 30

Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
35 40

<210> 1497
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1497
Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
20 25 30

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
35 40 45

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
50 55 60

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
65 70 75 80

Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
85 90 95

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
100 105 110

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
115 120 125

Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
130 135 140

<210> 1498
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1498
Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser
1 5 10 15

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr
20 25 30

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
35 40 45

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
50 55 60

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
65 70 75 80

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
85 90 95

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser
100 105 110

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
115 120 125

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
130 135 140

<210> 1499
<211> 288
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1499
Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
1 5 10 15

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
20 25 30

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
35 40 45

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
50 55 60

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
65 70 75 80

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro

85 90 95

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
100 105 110

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
115 120 125

Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
130 135 140

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
145 150 155 160

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
165 170 175

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
180 185 190

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
195 200 205

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
210 215 220

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
225 230 235 240

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
245 250 255

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
260 265 270

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
275 280 285

<210> 1500
<211> 288
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1500
Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
1 5 10 15

Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala
20 25 30

```
Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
        35                  40                  45

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
    50                  55                  60

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
65                  70                  75                  80

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
                85                  90                  95

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
            100                 105                 110

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
        115                 120                 125

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
    130                 135                 140

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro
145                 150                 155                 160

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
                165                 170                 175

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
            180                 185                 190

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
        195                 200                 205

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala
    210                 215                 220

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
225                 230                 235                 240

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
                245                 250                 255

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
            260                 265                 270

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
        275                 280                 285


<210> 1501
<211> 504
<212> PRT
```

<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 1501
Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
1 5 10 15

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
20 25 30

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
35 40 45

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Asn Pro
50 55 60

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
65 70 75 80

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
85 90 95

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
100 105 110

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
115 120 125

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
130 135 140

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
145 150 155 160

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
165 170 175

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
180 185 190

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Asn Pro
195 200 205

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
210 215 220

Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
225 230 235 240

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro

245 250 255

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
260 265 270

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
275 280 285

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
290 295 300

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
305 310 315 320

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
325 330 335

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
340 345 350

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
355 360 365

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
370 375 380

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
385 390 395 400

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
405 410 415

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
420 425 430

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
435 440 445

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
450 455 460

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
465 470 475 480

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
485 490 495

Gly Thr Ser Ser Thr Gly Ser Pro
500

<210> 1502

<211> 576
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1502

```
Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Gly Gly
1               5                  10                  15

Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Ser Glu Ser Gly Ser Ser
            20                  25                  30

Glu Gly Gly Pro Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro
        35                  40                  45

Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Ser Glu
    50                  55                  60

Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Ser Glu Ser Gly Ser Ser
65                  70                  75                  80

Glu Gly Gly Pro Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser
                85                  90                  95

Gly Ser Glu Gly Ser Ser Gly Pro Gly Glu Ser Ser Gly Ser Ser Glu
            100                 105                 110

Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Ser Glu Ser Gly Ser Ser
        115                 120                 125

Glu Gly Gly Pro Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro
    130                 135                 140

Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Glu Ser Pro
145                 150                 155                 160

Gly Gly Ser Ser Gly Ser Glu Ser Gly Glu Ser Pro Gly Gly Ser Ser
                165                 170                 175

Gly Ser Glu Ser Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser
            180                 185                 190

Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Gly Gly
        195                 200                 205

Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Gly Gly Glu Pro Ser Glu
    210                 215                 220

Ser Gly Ser Ser Gly Ser Glu Gly Ser Ser Gly Pro Gly Glu Ser Ser
225                 230                 235                 240
```

```
Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser Gly Ser Gly Gly
               245                 250                 255

Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Gly Gly Glu Pro Ser Glu
            260                 265                 270

Ser Gly Ser Ser Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser
        275                 280                 285

Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro Gly Glu Ser Pro
    290                 295                 300

Gly Gly Ser Ser Gly Ser Glu Ser Gly Glu Ser Pro Gly Gly Ser Ser
305                 310                 315                 320

Gly Ser Glu Ser Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser
               325                 330                 335

Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser Gly Glu Ser Pro
            340                 345                 350

Gly Gly Ser Ser Gly Ser Glu Ser Gly Ser Ser Glu Ser Gly Ser Ser
        355                 360                 365

Glu Gly Gly Pro Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser
    370                 375                 380

Gly Ser Glu Gly Ser Ser Gly Pro Gly Glu Ser Ser Gly Ser Ser Glu
385                 390                 395                 400

Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Gly Gly Glu Pro Ser Glu
               405                 410                 415

Ser Gly Ser Ser Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro
            420                 425                 430

Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Glu Ser Pro
        435                 440                 445

Gly Gly Ser Ser Gly Ser Glu Ser Gly Glu Ser Pro Gly Gly Ser Ser
    450                 455                 460

Gly Ser Glu Ser Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro
465                 470                 475                 480

Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Ser Glu
               485                 490                 495

Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Gly Gly Glu Pro Ser Glu
            500                 505                 510
```

Ser Gly Ser Ser Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser
515 520 525

Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser Gly Ser Glu Gly
530 535 540

Ser Ser Gly Pro Gly Glu Ser Ser Gly Ser Ser Glu Ser Gly Ser Ser
545 550 555 560

Glu Gly Gly Pro Gly Ser Glu Gly Ser Ser Gly Pro Gly Glu Ser Ser
565 570 575

<210> 1503
<211> 576
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1503
Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
35 40 45

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
50 55 60

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
65 70 75 80

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
85 90 95

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
100 105 110

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
115 120 125

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
130 135 140

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
145 150 155 160

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu

```
                165                 170                 175

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            180                 185                 190

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
        195                 200                 205

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
    210                 215                 220

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
225                 230                 235                 240

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
                245                 250                 255

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
            260                 265                 270

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        275                 280                 285

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
    290                 295                 300

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
305                 310                 315                 320

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
                325                 330                 335

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
            340                 345                 350

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
        355                 360                 365

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    370                 375                 380

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
385                 390                 395                 400

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
                405                 410                 415

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            420                 425                 430

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
```

435 440 445

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
450 455 460

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
465 470 475 480

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
485 490 495

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
500 505 510

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
515 520 525

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala
530 535 540

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
545 550 555 560

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
565 570 575

<210> 1504
<211> 576
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1504
Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser
1 5 10 15

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
20 25 30

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
35 40 45

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
50 55 60

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
65 70 75 80

Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
85 90 95

```
Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
            100                 105                 110

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser
        115                 120                 125

Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
    130                 135                 140

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
145                 150                 155                 160

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr
                165                 170                 175

Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
            180                 185                 190

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
        195                 200                 205

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly
    210                 215                 220

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
225                 230                 235                 240

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
                245                 250                 255

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
            260                 265                 270

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
        275                 280                 285

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
    290                 295                 300

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser
305                 310                 315                 320

Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
                325                 330                 335

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser
            340                 345                 350

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
        355                 360                 365
```

Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
370 375 380

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
385 390 395 400

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
405 410 415

Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
420 425 430

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro
435 440 445

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
450 455 460

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
465 470 475 480

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser
485 490 495

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser
500 505 510

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
515 520 525

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
530 535 540

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
545 550 555 560

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
565 570 575

<210> 1505
<211> 540
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1505
Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser
1 5 10 15

Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser Glu Ser Pro Ser

```
            20                  25                  30

Gly Thr Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro
        35                  40                  45

Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Thr
    50                  55                  60

Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser Glu Ser Pro Ser
65                  70                  75                  80

Gly Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro
                85                  90                  95

Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser
            100                 105                 110

Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser
        115                 120                 125

Ser Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
    130                 135                 140

Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Pro
145                 150                 155                 160

Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser
                165                 170                 175

Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
            180                 185                 190

Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Thr
        195                 200                 205

Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser Glu Ser Pro Ser
    210                 215                 220

Gly Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
225                 230                 235                 240

Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser
                245                 250                 255

Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Thr Pro Glu Ser Gly
            260                 265                 270

Ser Ala Ser Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
        275                 280                 285

Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Thr
```

290 295 300

Pro Glu Ser Gly Ser Ala Ser Pro Gly Thr Ser Thr Pro Glu Ser Gly
305 310 315 320

Ser Ala Ser Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
325 330 335

Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser
340 345 350

Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu
355 360 365

Ser Pro Gly Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
370 375 380

Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser
385 390 395 400

Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser
405 410 415

Gly Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
420 425 430

Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser
435 440 445

Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly
450 455 460

Ser Ala Ser Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro
465 470 475 480

Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro
485 490 495

Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu
500 505 510

Ser Pro Gly Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
515 520 525

Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
530 535 540

<210> 1506
<211> 836
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1506

```
Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Ser Glu
1               5                  10                  15

Ser Gly Ser Ser Glu Gly Gly Pro Gly Glu Ser Pro Gly Gly Ser Ser
            20                  25                  30

Gly Ser Glu Ser Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser
        35                  40                  45

Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser Gly Glu Ser Pro
    50                  55                  60

Gly Gly Ser Ser Gly Ser Glu Ser Gly Ser Ser Glu Ser Gly Ser Ser
65                  70                  75                  80

Glu Gly Gly Pro Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro
                85                  90                  95

Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro Gly Glu Ser Pro
            100                 105                 110

Gly Gly Ser Ser Gly Ser Glu Ser Gly Glu Ser Pro Gly Gly Ser Ser
        115                 120                 125

Gly Ser Glu Ser Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser
    130                 135                 140

Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Ser Glu
145                 150                 155                 160

Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Ser Glu Ser Gly Ser Ser
                165                 170                 175

Glu Gly Gly Pro Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro
            180                 185                 190

Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Ser Glu
        195                 200                 205

Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Gly Gly Glu Pro Ser Glu
    210                 215                 220

Ser Gly Ser Ser Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser
225                 230                 235                 240

Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser Gly Ser Gly Gly
                245                 250                 255
```

```
Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Glu Gly Ser Ser Gly Pro
            260                 265                 270

Gly Glu Ser Ser Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro
        275                 280                 285

Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Glu Gly
    290                 295                 300

Ser Ser Gly Pro Gly Glu Ser Ser Gly Ser Ser Glu Ser Gly Ser Ser
305                 310                 315                 320

Glu Gly Gly Pro Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser
                325                 330                 335

Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser Gly Ser Gly Gly
            340                 345                 350

Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Gly Gly Glu Pro Ser Glu
        355                 360                 365

Ser Gly Ser Ser Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro
    370                 375                 380

Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Gly Gly
385                 390                 395                 400

Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Glu Gly Ser Ser Gly Pro
                405                 410                 415

Gly Glu Ser Ser Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser
            420                 425                 430

Gly Ser Glu Gly Ser Ser Gly Pro Gly Glu Ser Ser Gly Ser Glu Gly
        435                 440                 445

Ser Ser Gly Pro Gly Glu Ser Ser Gly Ser Gly Gly Glu Pro Ser Glu
    450                 455                 460

Ser Gly Ser Ser Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro
465                 470                 475                 480

Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro Gly Glu Ser Pro
                485                 490                 495

Gly Gly Ser Ser Gly Ser Glu Ser Gly Ser Gly Gly Glu Pro Ser Glu
            500                 505                 510

Ser Gly Ser Ser Gly Ser Glu Gly Ser Ser Gly Pro Gly Glu Ser Ser
        515                 520                 525
```

```
Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser Gly Ser Glu Gly
    530                 535                 540

Ser Ser Gly Pro Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro
545                 550                 555                 560

Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Glu Gly
                565                 570                 575

Ser Ser Gly Pro Gly Glu Ser Ser Gly Ser Glu Gly Ser Ser Gly Pro
            580                 585                 590

Gly Glu Ser Ser Gly Ser Glu Gly Ser Ser Gly Pro Gly Glu Ser Ser
        595                 600                 605

Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Gly Gly
    610                 615                 620

Glu Pro Ser Glu Ser Gly Ser Ser Gly Glu Ser Pro Gly Gly Ser Ser
625                 630                 635                 640

Gly Ser Glu Ser Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser
                645                 650                 655

Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Glu Gly
            660                 665                 670

Ser Ser Gly Pro Gly Glu Ser Ser Gly Glu Ser Pro Gly Gly Ser Ser
        675                 680                 685

Gly Ser Glu Ser Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro
    690                 695                 700

Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Ser Glu
705                 710                 715                 720

Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Gly Gly Glu Pro Ser Glu
                725                 730                 735

Ser Gly Ser Ser Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro
            740                 745                 750

Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser Gly Ser Gly Gly
        755                 760                 765

Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Ser Glu Ser Gly Ser Ser
    770                 775                 780

Glu Gly Gly Pro Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser
785                 790                 795                 800
```

Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Glu Ser Pro
805 810 815

Gly Gly Ser Ser Gly Ser Glu Ser Gly Ser Gly Gly Glu Pro Ser Glu
820 825 830

Ser Gly Ser Ser
835

<210> 1507
<211> 864
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1507
Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
35 40 45

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
50 55 60

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
65 70 75 80

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
85 90 95

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
100 105 110

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
115 120 125

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
130 135 140

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
145 150 155 160

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
165 170 175

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro

180 185 190

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
195 200 205

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
210 215 220

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
225 230 235 240

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
245 250 255

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
260 265 270

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
275 280 285

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
290 295 300

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
305 310 315 320

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
325 330 335

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
340 345 350

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
355 360 365

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
370 375 380

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
385 390 395 400

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
405 410 415

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
420 425 430

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
435 440 445

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro

```
    450                 455                 460

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
465                 470                 475                 480

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
                485                 490                 495

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
            500                 505                 510

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
        515                 520                 525

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala
    530                 535                 540

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
545                 550                 555                 560

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
                565                 570                 575

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
            580                 585                 590

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
        595                 600                 605

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
    610                 615                 620

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
625                 630                 635                 640

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
                645                 650                 655

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
            660                 665                 670

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
        675                 680                 685

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
    690                 695                 700

Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
705                 710                 715                 720

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
```

725 730 735

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
740 745 750

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
755 760 765

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
770 775 780

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
785 790 795 800

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
805 810 815

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
820 825 830

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
835 840 845

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
850 855 860

<210> 1508
<211> 875
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<220>
<221> MOD_RES
<222> (430)..(432)
<223> Any amino acid

<220>
<221> MOD_RES
<222> (443)..(446)
<223> Any amino acid

<400> 1508
Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Pro
1 5 10 15

Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser
20 25 30

Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
35 40 45

```
Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Thr Ser Thr
    50                  55                  60

Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser Glu Ser Pro Ser
65                  70                  75                  80

Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
                85                  90                  95

Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser
            100                 105                 110

Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser
        115                 120                 125

Ser Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro
    130                 135                 140

Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro
145                 150                 155                 160

Ser Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser
                165                 170                 175

Ser Thr Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro
            180                 185                 190

Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Thr Ser Thr
        195                 200                 205

Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser Glu Ser Pro Ser
    210                 215                 220

Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
225                 230                 235                 240

Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser
                245                 250                 255

Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Thr Pro Glu Ser Gly
            260                 265                 270

Ser Ala Ser Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
        275                 280                 285

Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser
    290                 295                 300

Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro Ser Gly Glu Ser
305                 310                 315                 320
```

```
Ser Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
                325                 330                 335

Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser
            340                 345                 350

Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser Ser Thr Ala Glu
        355                 360                 365

Ser Pro Gly Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro
    370                 375                 380

Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser
385                 390                 395                 400

Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser
                405                 410                 415

Gly Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Pro Xaa Xaa Xaa
            420                 425                 430

Gly Ala Ser Ala Ser Gly Ala Pro Ser Thr Xaa Xaa Xaa Xaa Ser Glu
        435                 440                 445

Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly
    450                 455                 460

Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly
465                 470                 475                 480

Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu
                485                 490                 495

Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly
            500                 505                 510

Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly
        515                 520                 525

Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Pro Ser
    530                 535                 540

Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser
545                 550                 555                 560

Pro Gly Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly
                565                 570                 575

Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser Glu
            580                 585                 590
```

```
Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly
        595                 600                 605

Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly
    610                 615                 620

Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Thr Pro
625                 630                 635                 640

Glu Ser Gly Ser Ala Ser Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser
                645                 650                 655

Ala Ser Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly
            660                 665                 670

Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser Ser
        675                 680                 685

Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly
    690                 695                 700

Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly
705                 710                 715                 720

Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser Ser
                725                 730                 735

Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser
            740                 745                 750

Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly
        755                 760                 765

Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Pro Ser
    770                 775                 780

Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser
785                 790                 795                 800

Thr Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly
                805                 810                 815

Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro Ser
            820                 825                 830

Gly Glu Ser Ser Thr Ala Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
        835                 840                 845

Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
    850                 855                 860
```

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
865 870 875

<210> 1509
<211> 864
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1509
Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
1 5 10 15

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
20 25 30

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
35 40 45

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
50 55 60

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
65 70 75 80

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
85 90 95

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
100 105 110

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
115 120 125

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
130 135 140

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
145 150 155 160

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
165 170 175

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
180 185 190

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
195 200 205

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
210 215 220

```
Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
225                 230                 235                 240

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro
                245                 250                 255

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
            260                 265                 270

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
        275                 280                 285

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
    290                 295                 300

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
305                 310                 315                 320

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                325                 330                 335

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
            340                 345                 350

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
        355                 360                 365

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    370                 375                 380

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
385                 390                 395                 400

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
                405                 410                 415

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
            420                 425                 430

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
        435                 440                 445

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
    450                 455                 460

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
465                 470                 475                 480

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
                485                 490                 495
```

```
Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
            500                 505                 510

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
        515                 520                 525

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
    530                 535                 540

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
545                 550                 555                 560

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                565                 570                 575

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
            580                 585                 590

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
        595                 600                 605

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
    610                 615                 620

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
625                 630                 635                 640

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
                645                 650                 655

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            660                 665                 670

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
        675                 680                 685

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
    690                 695                 700

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
705                 710                 715                 720

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
                725                 730                 735

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
            740                 745                 750

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
        755                 760                 765
```

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
770 775 780

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
785 790 795 800

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
805 810 815

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
820 825 830

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
835 840 845

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
850 855 860

<210> 1510
<211> 875
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1510
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
1 5 10 15

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
20 25 30

Ser Thr Glu Glu Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro
35 40 45

Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser
50 55 60

Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser
65 70 75 80

Gly Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
85 90 95

Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Glu Pro
100 105 110

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
115 120 125

```
Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
    130                 135                 140

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
145                 150                 155                 160

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
                165                 170                 175

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            180                 185                 190

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
        195                 200                 205

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
    210                 215                 220

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
225                 230                 235                 240

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
                245                 250                 255

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
            260                 265                 270

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        275                 280                 285

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
    290                 295                 300

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
305                 310                 315                 320

Ser Thr Glu Glu Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
                325                 330                 335

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
            340                 345                 350

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Ser Thr Glu Pro Ser Glu
        355                 360                 365

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    370                 375                 380

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
385                 390                 395                 400
```

```
Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
                405                 410                 415

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            420                 425                 430

Gly Ala Ser Ala Ser Gly Ala Pro Ser Thr Gly Gly Thr Ser Glu Ser
        435                 440                 445

Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
    450                 455                 460

Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
465                 470                 475                 480

Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser Glu
                485                 490                 495

Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser
            500                 505                 510

Thr Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
        515                 520                 525

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser
    530                 535                 540

Ala Ser Thr Gly Thr Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
545                 550                 555                 560

Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
                565                 570                 575

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Thr Ser Ser
            580                 585                 590

Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser
        595                 600                 605

Pro Gly Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly
    610                 615                 620

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala
625                 630                 635                 640

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                645                 650                 655

Ser Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly
            660                 665                 670
```

Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser Glu
675 680 685

Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
690 695 700

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
705 710 715 720

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Ser Thr Pro
725 730 735

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
740 745 750

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
755 760 765

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
770 775 780

Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
785 790 795 800

Thr Glu Glu Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
805 810 815

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly
820 825 830

Thr Ser Ser Thr Gly Ser Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
835 840 845

Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
850 855 860

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
865 870 875

<210> 1511
<211> 1318
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1511
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
1 5 10 15

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
20 25 30

```
Ser Thr Glu Glu Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro
        35                  40                  45

Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser
    50                  55                  60

Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser
65                  70                  75                  80

Gly Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
                85                  90                  95

Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Glu Pro
            100                 105                 110

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
        115                 120                 125

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
    130                 135                 140

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
145                 150                 155                 160

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
                165                 170                 175

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            180                 185                 190

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
        195                 200                 205

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
    210                 215                 220

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
225                 230                 235                 240

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
                245                 250                 255

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
            260                 265                 270

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        275                 280                 285

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
    290                 295                 300
```

```
Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
305                 310                 315                 320

Ser Thr Glu Glu Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
                325                 330                 335

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
            340                 345                 350

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Ser Thr Glu Pro Ser Glu
        355                 360                 365

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    370                 375                 380

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
385                 390                 395                 400

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
                405                 410                 415

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            420                 425                 430

Gly Pro Glu Pro Thr Gly Pro Ala Pro Ser Gly Gly Ser Glu Pro Ala
        435                 440                 445

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
    450                 455                 460

Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
465                 470                 475                 480

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly
                485                 490                 495

Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser
            500                 505                 510

Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
        515                 520                 525

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
    530                 535                 540

Ser Pro Thr Ser Thr Glu Glu Gly Ser Thr Ser Ser Thr Ala Glu Ser
545                 550                 555                 560

Pro Gly Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly
                565                 570                 575
```

```
Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser Glu
            580                 585                 590

Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly
        595                 600                 605

Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly
    610                 615                 620

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
625                 630                 635                 640

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
                645                 650                 655

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
            660                 665                 670

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser
        675                 680                 685

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    690                 695                 700

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
705                 710                 715                 720

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Pro Ser
                725                 730                 735

Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser
            740                 745                 750

Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly
        755                 760                 765

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
    770                 775                 780

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
785                 790                 795                 800

Ser Ala Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
                805                 810                 815

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
            820                 825                 830

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
        835                 840                 845
```

```
Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
    850                 855                 860

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Ala Ser
865                 870                 875                 880

Gly Ala Pro Ser Thr Gly Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr
                885                 890                 895

Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr
            900                 905                 910

Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Glu Ser Ala
        915                 920                 925

Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
    930                 935                 940

Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
945                 950                 955                 960

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser
                965                 970                 975

Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
            980                 985                 990

Ser Pro Gly Thr Ser Thr Pro Glu  Ser Gly Ser Ala Ser  Pro Gly Thr
        995                 1000                1005

Ser Pro  Ser Gly Glu Ser Ser  Thr Ala Pro Gly Thr  Ser Pro Ser
    1010                1015                1020

Gly Glu  Ser Ser Thr Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1025                1030                1035

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1040                1045                1050

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    1055                1060                1065

Thr Ser  Glu Ser Pro Ser Gly  Thr Ala Pro Gly Ser  Thr Ser Glu
    1070                1075                1080

Ser Pro  Ser Gly Thr Ala Pro  Gly Thr Ser Thr Pro  Glu Ser Gly
    1085                1090                1095

Ser Ala  Ser Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    1100                1105                1110
```

Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1115 1120 1125

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
1130 1135 1140

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
1145 1150 1155

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1160 1165 1170

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala
1175 1180 1185

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro
1190 1195 1200

Ser Gly Ala Thr Gly Ser Pro Gly Ser Thr Ser Glu Ser Pro Ser
1205 1210 1215

Gly Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala
1220 1225 1230

Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser
1235 1240 1245

Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly
1250 1255 1260

Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
1265 1270 1275

Ser Ser Ser Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
1280 1285 1290

Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr
1295 1300 1305

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
1310 1315

<210> 1512
<211> 924
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1512

```
Met Ala Glu Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ala Ser
1               5                   10                  15

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly
            20                  25                  30

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
        35                  40                  45

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu
    50                  55                  60

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro
65                  70                  75                  80

Thr Ser Thr Glu Glu Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly
                85                  90                  95

Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr
            100                 105                 110

Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro
        115                 120                 125

Ser Gly Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser
    130                 135                 140

Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Glu
145                 150                 155                 160

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
                165                 170                 175

Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
            180                 185                 190

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
        195                 200                 205

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser
    210                 215                 220

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
225                 230                 235                 240

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
                245                 250                 255

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
            260                 265                 270
```

```
Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
        275                 280                 285

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
    290                 295                 300

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
305                 310                 315                 320

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
                325                 330                 335

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu
            340                 345                 350

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro
        355                 360                 365

Thr Ser Thr Glu Glu Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
    370                 375                 380

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser
385                 390                 395                 400

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Ser Thr Glu Pro Ser
                405                 410                 415

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            420                 425                 430

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro
        435                 440                 445

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro
    450                 455                 460

Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
465                 470                 475                 480

Pro Gly Ala Ser Ala Ser Gly Ala Pro Ser Thr Gly Gly Thr Ser Glu
                485                 490                 495

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
            500                 505                 510

Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
        515                 520                 525

Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser
    530                 535                 540
```

```
Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser
545                 550                 555                 560

Ser Thr Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
                565                 570                 575

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
            580                 585                 590

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
        595                 600                 605

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
    610                 615                 620

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Thr Ser
625                 630                 635                 640

Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser Ser Thr Ala Glu
                645                 650                 655

Ser Pro Gly Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro
            660                 665                 670

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
        675                 680                 685

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu
    690                 695                 700

Gly Ser Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro
705                 710                 715                 720

Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser
                725                 730                 735

Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
            740                 745                 750

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
        755                 760                 765

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Ser Thr
    770                 775                 780

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
785                 790                 795                 800

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                805                 810                 815
```

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
820 825 830

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
835 840 845

Ser Thr Glu Glu Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
850 855 860

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
865 870 875 880

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Ser Glu Ser Ala Thr Pro
885 890 895

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
900 905 910

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
915 920

<210> 1513
<211> 913
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1513
Met Ala Glu Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Pro
1 5 10 15

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
20 25 30

Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
35 40 45

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
50 55 60

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser
65 70 75 80

Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
85 90 95

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
100 105 110

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
115 120 125

```
Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
    130                 135                 140

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro
145                 150                 155                 160

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr
                165                 170                 175

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            180                 185                 190

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro
        195                 200                 205

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
    210                 215                 220

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
225                 230                 235                 240

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
                245                 250                 255

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
            260                 265                 270

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
        275                 280                 285

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
    290                 295                 300

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
305                 310                 315                 320

Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
                325                 330                 335

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            340                 345                 350

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        355                 360                 365

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    370                 375                 380

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
385                 390                 395                 400
```

```
Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                405                 410                 415

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            420                 425                 430

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
        435                 440                 445

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro
    450                 455                 460

Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
465                 470                 475                 480

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
                485                 490                 495

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
            500                 505                 510

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
        515                 520                 525

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
    530                 535                 540

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
545                 550                 555                 560

Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
                565                 570                 575

Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro
            580                 585                 590

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr
        595                 600                 605

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
    610                 615                 620

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
625                 630                 635                 640

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
                645                 650                 655

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
            660                 665                 670
```

```
Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
        675                 680                 685

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro
    690                 695                 700

Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
705                 710                 715                 720

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser
                725                 730                 735

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro
            740                 745                 750

Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
        755                 760                 765

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
    770                 775                 780

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
785                 790                 795                 800

Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
                805                 810                 815

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu
            820                 825                 830

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro
        835                 840                 845

Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
    850                 855                 860

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu
865                 870                 875                 880

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
                885                 890                 895

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            900                 905                 910

Pro
```

<210> 1514
<211> 864

<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1514

```
Gly Thr Ser Thr Glu Pro Ser Glu Pro Gly Ser Ala Gly Thr Ser Thr
1               5                  10                  15

Glu Pro Ser Glu Pro Gly Ser Ala Gly Ser Glu Pro Ala Thr Ser Gly
            20                  25                  30

Thr Glu Pro Ser Gly Ser Gly Ala Ser Glu Pro Thr Ser Thr Glu Pro
        35                  40                  45

Gly Ser Glu Pro Ala Thr Ser Gly Thr Glu Pro Ser Gly Ser Glu Pro
    50                  55                  60

Ala Thr Ser Gly Thr Glu Pro Ser Gly Ser Glu Pro Ala Thr Ser Gly
65                  70                  75                  80

Thr Glu Pro Ser Gly Ser Gly Ala Ser Glu Pro Thr Ser Thr Glu Pro
                85                  90                  95

Gly Thr Ser Thr Glu Pro Ser Glu Pro Gly Ser Ala Gly Ser Glu Pro
            100                 105                 110

Ala Thr Ser Gly Thr Glu Pro Ser Gly Thr Ser Thr Glu Pro Ser Glu
        115                 120                 125

Pro Gly Ser Ala Gly Ser Glu Pro Ala Thr Ser Gly Thr Glu Pro Ser
    130                 135                 140

Gly Ser Glu Pro Ala Thr Ser Gly Thr Glu Pro Ser Gly Thr Ser Thr
145                 150                 155                 160

Glu Pro Ser Glu Pro Gly Ser Ala Gly Thr Ser Thr Glu Pro Ser Glu
                165                 170                 175

Pro Gly Ser Ala Gly Ser Glu Pro Ala Thr Ser Gly Thr Glu Pro Ser
            180                 185                 190

Gly Ser Glu Pro Ala Thr Ser Gly Thr Glu Pro Ser Gly Thr Ser Glu
        195                 200                 205

Pro Ser Thr Ser Glu Pro Gly Ala Gly Ser Gly Ala Ser Glu Pro Thr
    210                 215                 220

Ser Thr Glu Pro Gly Thr Ser Glu Pro Ser Thr Ser Glu Pro Gly Ala
225                 230                 235                 240
```

```
Gly Ser Glu Pro Ala Thr Ser Gly Thr Glu Pro Ser Gly Ser Glu Pro
                245                 250                 255

Ala Thr Ser Gly Thr Glu Pro Ser Gly Thr Ser Thr Glu Pro Ser Glu
            260                 265                 270

Pro Gly Ser Ala Gly Thr Ser Thr Glu Pro Ser Glu Pro Gly Ser Ala
        275                 280                 285

Gly Ser Gly Ala Ser Glu Pro Thr Ser Thr Glu Pro Gly Ser Glu Pro
    290                 295                 300

Ala Thr Ser Gly Thr Glu Pro Ser Gly Ser Glu Pro Ala Thr Ser Gly
305                 310                 315                 320

Thr Glu Pro Ser Gly Ser Glu Pro Ala Thr Ser Gly Thr Glu Pro Ser
                325                 330                 335

Gly Ser Glu Pro Ala Thr Ser Gly Thr Glu Pro Ser Gly Thr Ser Thr
            340                 345                 350

Glu Pro Ser Glu Pro Gly Ser Ala Gly Ser Glu Pro Ala Thr Ser Gly
        355                 360                 365

Thr Glu Pro Ser Gly Ser Gly Ala Ser Glu Pro Thr Ser Thr Glu Pro
    370                 375                 380

Gly Thr Ser Thr Glu Pro Ser Glu Pro Gly Ser Ala Gly Ser Glu Pro
385                 390                 395                 400

Ala Thr Ser Gly Thr Glu Pro Ser Gly Ser Gly Ala Ser Glu Pro Thr
                405                 410                 415

Ser Thr Glu Pro Gly Thr Ser Thr Glu Pro Ser Glu Pro Gly Ser Ala
            420                 425                 430

Gly Ser Gly Ala Ser Glu Pro Thr Ser Thr Glu Pro Gly Ser Glu Pro
        435                 440                 445

Ala Thr Ser Gly Thr Glu Pro Ser Gly Ser Gly Ala Ser Glu Pro Thr
    450                 455                 460

Ser Thr Glu Pro Gly Ser Glu Pro Ala Thr Ser Gly Thr Glu Pro Ser
465                 470                 475                 480

Gly Ser Gly Ala Ser Glu Pro Thr Ser Thr Glu Pro Gly Thr Ser Thr
                485                 490                 495

Glu Pro Ser Glu Pro Gly Ser Ala Gly Ser Glu Pro Ala Thr Ser Gly
            500                 505                 510
```

```
Thr Glu Pro Ser Gly Ser Gly Ala Ser Glu Pro Thr Ser Thr Glu Pro
        515                 520                 525

Gly Thr Ser Thr Glu Pro Ser Glu Pro Gly Ser Ala Gly Ser Glu Pro
    530                 535                 540

Ala Thr Ser Gly Thr Glu Pro Ser Gly Thr Ser Thr Glu Pro Ser Glu
545                 550                 555                 560

Pro Gly Ser Ala Gly Ser Glu Pro Ala Thr Ser Gly Thr Glu Pro Ser
                565                 570                 575

Gly Thr Ser Thr Glu Pro Ser Glu Pro Gly Ser Ala Gly Thr Ser Thr
            580                 585                 590

Glu Pro Ser Glu Pro Gly Ser Ala Gly Thr Ser Thr Glu Pro Ser Glu
        595                 600                 605

Pro Gly Ser Ala Gly Thr Ser Thr Glu Pro Ser Glu Pro Gly Ser Ala
    610                 615                 620

Gly Thr Ser Thr Glu Pro Ser Glu Pro Gly Ser Ala Gly Thr Ser Thr
625                 630                 635                 640

Glu Pro Ser Glu Pro Gly Ser Ala Gly Thr Ser Glu Pro Ser Thr Ser
                645                 650                 655

Glu Pro Gly Ala Gly Ser Gly Ala Ser Glu Pro Thr Ser Thr Glu Pro
            660                 665                 670

Gly Thr Ser Thr Glu Pro Ser Glu Pro Gly Ser Ala Gly Thr Ser Thr
        675                 680                 685

Glu Pro Ser Glu Pro Gly Ser Ala Gly Thr Ser Thr Glu Pro Ser Glu
    690                 695                 700

Pro Gly Ser Ala Gly Ser Glu Pro Ala Thr Ser Gly Thr Glu Pro Ser
705                 710                 715                 720

Gly Ser Gly Ala Ser Glu Pro Thr Ser Thr Glu Pro Gly Ser Glu Pro
                725                 730                 735

Ala Thr Ser Gly Thr Glu Pro Ser Gly Ser Glu Pro Ala Thr Ser Gly
            740                 745                 750

Thr Glu Pro Ser Gly Ser Glu Pro Ala Thr Ser Gly Thr Glu Pro Ser
        755                 760                 765

Gly Ser Glu Pro Ala Thr Ser Gly Thr Glu Pro Ser Gly Thr Ser Glu
    770                 775                 780
```

Pro Ser Thr Ser Glu Pro Gly Ala Gly Ser Glu Pro Ala Thr Ser Gly
785 790 795 800

Thr Glu Pro Ser Gly Ser Gly Ala Ser Glu Pro Thr Ser Thr Glu Pro
805 810 815

Gly Thr Ser Thr Glu Pro Ser Glu Pro Gly Ser Ala Gly Ser Glu Pro
820 825 830

Ala Thr Ser Gly Thr Glu Pro Ser Gly Ser Gly Ala Ser Glu Pro Thr
835 840 845

Ser Thr Glu Pro Gly Thr Ser Thr Glu Pro Ser Glu Pro Gly Ser Ala
850 855 860

<210> 1515
<211> 288
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1515
Gly Ser Gly Gly Glu Gly Gly Ser Gly Gly Glu Gly Gly Ser Gly Gly
1 5 10 15

Glu Gly Gly Ser Gly Gly Glu Gly Gly Ser Gly Gly Glu Gly Gly Ser
20 25 30

Gly Gly Glu Gly Gly Ser Gly Gly Glu Gly Gly Ser Gly Gly Glu Gly
35 40 45

Gly Ser Gly Gly Glu Gly Gly Ser Gly Gly Glu Gly Gly Ser Gly Gly
50 55 60

Glu Gly Gly Ser Gly Gly Glu Gly Gly Ser Gly Gly Glu Gly Gly Ser
65 70 75 80

Gly Gly Glu Gly Gly Ser Gly Gly Glu Gly Gly Ser Gly Gly Glu Gly
85 90 95

Gly Ser Gly Gly Glu Gly Gly Ser Gly Gly Glu Gly Gly Ser Gly Gly
100 105 110

Glu Gly Gly Ser Gly Gly Glu Gly Gly Ser Gly Gly Glu Gly Gly Ser
115 120 125

Gly Gly Glu Gly Gly Ser Gly Gly Glu Gly Gly Ser Gly Gly Glu Gly
130 135 140

Gly Ser Gly Gly Glu Gly Gly Ser Gly Gly Glu Gly Gly Ser Gly Gly
145 150 155 160

Glu Gly Gly Ser Gly Gly Glu Gly Gly Ser Gly Gly Glu Gly Gly Ser
165 170 175

Gly Gly Glu Gly Gly Ser Gly Gly Glu Gly Gly Ser Gly Gly Glu Gly
180 185 190

Gly Ser Gly Gly Glu Gly Gly Ser Gly Gly Glu Gly Gly Ser Gly Gly
195 200 205

Glu Gly Gly Ser Gly Gly Glu Gly Gly Ser Gly Gly Glu Gly Gly Ser
210 215 220

Gly Gly Glu Gly Gly Ser Gly Gly Glu Gly Gly Ser Gly Gly Glu Gly
225 230 235 240

Gly Ser Gly Gly Glu Gly Gly Ser Gly Gly Glu Gly Gly Ser Gly Gly
245 250 255

Glu Gly Gly Ser Gly Gly Glu Gly Gly Ser Gly Gly Glu Gly Gly Ser
260 265 270

Gly Gly Glu Gly Gly Ser Gly Gly Glu Gly Gly Ser Gly Gly Glu Gly
275 280 285

<210> 1516
<211> 288
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1516
Gly Glu Gly Glu Gly Gly Gly Glu Gly Gly Glu Gly Glu Gly Gly Gly
1 5 10 15

Glu Gly Gly Glu Gly Glu Gly Gly Gly Glu Gly Gly Glu Gly Glu Gly
20 25 30

Gly Gly Glu Gly Gly Glu Gly Glu Gly Gly Gly Glu Gly Gly Glu Gly
35 40 45

Glu Gly Gly Gly Glu Gly Gly Glu Gly Glu Gly Gly Gly Glu Gly Gly
50 55 60

Glu Gly Glu Gly Gly Gly Glu Gly Gly Glu Gly Glu Gly Gly Gly Glu
65 70 75 80

Gly Gly Glu Gly Glu Gly Gly Gly Glu Gly Gly Glu Gly Glu Gly Gly
85 90 95

Gly Glu Gly Gly Glu Gly Glu Gly Gly Gly Glu Gly Gly Glu Gly Glu
100 105 110

Gly Gly Gly Glu Gly Gly Glu Gly Glu Gly Gly Gly Glu Gly Gly Glu
115 120 125

Gly Glu Gly Gly Gly Glu Gly Gly Glu Gly Glu Gly Gly Gly Glu Gly
130 135 140

Gly Glu Gly Glu Gly Gly Gly Glu Gly Gly Glu Gly Glu Gly Gly Gly
145 150 155 160

Glu Gly Gly Glu Gly Glu Gly Gly Gly Glu Gly Gly Glu Gly Glu Gly
165 170 175

Gly Gly Glu Gly Gly Glu Gly Glu Gly Gly Gly Glu Gly Gly Glu Gly
180 185 190

Glu Gly Gly Gly Glu Gly Gly Glu Gly Glu Gly Gly Gly Glu Gly Gly
195 200 205

Glu Gly Glu Gly Gly Gly Glu Gly Gly Glu Gly Glu Gly Gly Gly Glu
210 215 220

Gly Gly Glu Gly Glu Gly Gly Gly Glu Gly Gly Glu Gly Glu Gly Gly
225 230 235 240

Gly Glu Gly Gly Glu Gly Glu Gly Gly Gly Glu Gly Gly Glu Gly Glu
245 250 255

Gly Gly Gly Glu Gly Gly Glu Gly Glu Gly Gly Gly Glu Gly Gly Glu
260 265 270

Gly Glu Gly Gly Gly Glu Gly Gly Glu Gly Glu Gly Gly Gly Glu Gly
275 280 285

<210> 1517
<211> 288
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1517
Ser Ser Glu Ser Ser Glu Ser Ser Ser Ser Glu Ser Ser Ser Glu Ser
1 5 10 15

Ser Glu Ser Ser Ser Ser Glu Ser Ser Ser Glu Ser Ser Glu Ser Ser
20 25 30

Ser Ser Glu Ser Ser Ser Glu Ser Ser Glu Ser Ser Ser Ser Glu Ser
35 40 45

Ser Ser Glu Ser Ser Glu Ser Ser Ser Ser Glu Ser Ser Ser Glu Ser
50 55 60

Ser Glu Ser Ser Ser Ser Glu Ser Ser Ser Glu Ser Ser Glu Ser Ser
65 70 75 80

Ser Ser Glu Ser Ser Ser Glu Ser Ser Glu Ser Ser Ser Ser Glu Ser
85 90 95

Ser Ser Glu Ser Ser Glu Ser Ser Ser Ser Glu Ser Ser Ser Glu Ser
100 105 110

Ser Glu Ser Ser Ser Ser Glu Ser Ser Ser Glu Ser Ser Glu Ser Ser
115 120 125

Ser Ser Glu Ser Ser Ser Glu Ser Ser Glu Ser Ser Ser Ser Glu Ser
130 135 140

Ser Ser Glu Ser Ser Glu Ser Ser Ser Ser Glu Ser Ser Ser Glu Ser
145 150 155 160

Ser Glu Ser Ser Ser Ser Glu Ser Ser Ser Glu Ser Ser Glu Ser Ser
165 170 175

Ser Ser Glu Ser Ser Ser Glu Ser Ser Glu Ser Ser Ser Ser Glu Ser
180 185 190

Ser Ser Glu Ser Ser Glu Ser Ser Ser Ser Glu Ser Ser Ser Glu Ser
195 200 205

Ser Glu Ser Ser Ser Ser Glu Ser Ser Ser Glu Ser Ser Glu Ser Ser
210 215 220

Ser Ser Glu Ser Ser Ser Glu Ser Ser Glu Ser Ser Ser Ser Glu Ser
225 230 235 240

Ser Ser Glu Ser Ser Glu Ser Ser Ser Ser Glu Ser Ser Ser Glu Ser
245 250 255

Ser Glu Ser Ser Ser Ser Glu Ser Ser Ser Glu Ser Ser Glu Ser Ser
260 265 270

Ser Ser Glu Ser Ser Ser Glu Ser Ser Glu Ser Ser Ser Ser Glu Ser
275 280 285

<210> 1518
<211> 288
<212> PRT
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1518
Gly Glu Gly Ser Gly Glu Gly Ser Glu Gly Glu Gly Ser Glu Gly Ser
1 5 10 15

Gly Glu Gly Glu Gly Ser Glu Gly Ser Gly Glu Gly Glu Gly Gly Ser
20 25 30

Glu Gly Ser Glu Gly Glu Gly Gly Ser Glu Gly Ser Glu Gly Glu Gly
35 40 45

Gly Ser Glu Gly Ser Glu Gly Glu Gly Ser Gly Glu Gly Ser Glu Gly
50 55 60

Glu Gly Gly Ser Glu Gly Ser Glu Gly Glu Gly Ser Gly Glu Gly Ser
65 70 75 80

Glu Gly Glu Gly Ser Glu Gly Gly Ser Glu Gly Glu Gly Gly Ser Glu
85 90 95

Gly Ser Glu Gly Glu Gly Ser Gly Glu Gly Ser Glu Gly Glu Gly Gly
100 105 110

Glu Gly Gly Ser Glu Gly Glu Gly Ser Glu Gly Ser Gly Glu Gly Glu
115 120 125

Gly Ser Gly Glu Gly Ser Glu Gly Glu Gly Ser Glu Gly Ser Gly Glu
130 135 140

Gly Glu Gly Ser Gly Glu Gly Ser Glu Gly Glu Gly Ser Glu Gly Ser
145 150 155 160

Gly Glu Gly Glu Gly Ser Glu Gly Ser Gly Glu Gly Glu Gly Gly Ser
165 170 175

Glu Gly Ser Glu Gly Glu Gly Ser Glu Gly Ser Gly Glu Gly Glu Gly
180 185 190

Gly Glu Gly Ser Gly Glu Gly Glu Gly Ser Gly Glu Gly Ser Glu Gly
195 200 205

Glu Gly Gly Gly Glu Gly Ser Glu Gly Glu Gly Ser Gly Glu Gly Gly
210 215 220

Glu Gly Glu Gly Ser Glu Gly Gly Ser Glu Gly Glu Gly Gly Ser Glu
225 230 235 240

Gly Gly Glu Gly Glu Gly Ser Glu Gly Ser Gly Glu Gly Glu Gly Ser
245 250 255

Glu Gly Gly Ser Glu Gly Glu Gly Ser Glu Gly Gly Ser Glu Gly Glu
260 265 270

Gly Ser Glu Gly Ser Gly Glu Gly Glu Gly Ser Glu Gly Ser Gly Glu
275 280 285

<210> 1519
<211> 576
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1519
Gly Gly Lys Pro Gly Glu Gly Gly Lys Pro Glu Gly Gly Gly Gly Lys
1 5 10 15

Pro Gly Gly Lys Pro Glu Gly Glu Gly Glu Gly Lys Pro Gly Gly Lys
20 25 30

Pro Glu Gly Gly Gly Lys Pro Gly Gly Gly Glu Gly Gly Lys Pro Glu
35 40 45

Gly Gly Lys Pro Glu Gly Glu Gly Lys Pro Gly Gly Gly Glu Gly Lys
50 55 60

Pro Gly Gly Lys Pro Glu Gly Gly Gly Gly Lys Pro Glu Gly Glu Gly
65 70 75 80

Lys Pro Gly Gly Gly Gly Gly Lys Pro Gly Gly Lys Pro Glu Gly Glu
85 90 95

Gly Lys Pro Gly Gly Gly Glu Gly Gly Lys Pro Glu Gly Lys Pro Gly
100 105 110

Glu Gly Gly Glu Gly Lys Pro Gly Gly Lys Pro Glu Gly Gly Gly Glu
115 120 125

Gly Lys Pro Gly Gly Gly Lys Pro Gly Glu Gly Gly Lys Pro Gly Glu
130 135 140

Gly Lys Pro Gly Gly Gly Glu Gly Gly Lys Pro Glu Gly Gly Lys Pro
145 150 155 160

Glu Gly Glu Gly Lys Pro Gly Gly Gly Glu Gly Lys Pro Gly Gly Lys
165 170 175

Pro Gly Glu Gly Gly Lys Pro Glu Gly Gly Gly Glu Gly Lys Pro Gly
180 185 190

Gly Lys Pro Gly Glu Gly Gly Glu Gly Lys Pro Gly Gly Gly Lys Pro
195 200 205

```
Glu Gly Glu Gly Lys Pro Gly Gly Gly Lys Pro Gly Gly Gly Glu Gly
    210                 215                 220

Gly Lys Pro Glu Gly Glu Gly Lys Pro Gly Gly Lys Pro Glu Gly Gly
225                 230                 235                 240

Gly Glu Gly Lys Pro Gly Gly Lys Pro Glu Gly Gly Gly Lys Pro Glu
                245                 250                 255

Gly Gly Gly Glu Gly Lys Pro Gly Gly Gly Lys Pro Gly Glu Gly Gly
            260                 265                 270

Lys Pro Gly Glu Gly Glu Gly Lys Pro Gly Gly Lys Pro Glu Gly Glu
        275                 280                 285

Gly Lys Pro Gly Gly Glu Gly Gly Gly Lys Pro Glu Gly Lys Pro Gly
    290                 295                 300

Gly Gly Glu Gly Gly Lys Pro Glu Gly Gly Lys Pro Gly Glu Gly Gly
305                 310                 315                 320

Lys Pro Glu Gly Gly Lys Pro Gly Glu Gly Gly Glu Gly Lys Pro Gly
                325                 330                 335

Gly Gly Lys Pro Gly Glu Gly Gly Lys Pro Glu Gly Gly Gly Lys Pro
            340                 345                 350

Glu Gly Glu Gly Lys Pro Gly Gly Gly Gly Lys Pro Gly Glu Gly Gly
        355                 360                 365

Lys Pro Glu Gly Gly Lys Pro Glu Gly Gly Gly Glu Gly Lys Pro Gly
    370                 375                 380

Gly Gly Lys Pro Glu Gly Glu Gly Lys Pro Gly Gly Gly Glu Gly Lys
385                 390                 395                 400

Pro Gly Gly Lys Pro Glu Gly Gly Gly Gly Lys Pro Gly Glu Gly Gly
                405                 410                 415

Lys Pro Glu Gly Gly Lys Pro Gly Gly Glu Gly Gly Gly Lys Pro Glu
            420                 425                 430

Gly Glu Gly Lys Pro Gly Gly Lys Pro Gly Glu Gly Gly Gly Gly Lys
        435                 440                 445

Pro Gly Gly Lys Pro Glu Gly Glu Gly Lys Pro Gly Glu Gly Gly Glu
    450                 455                 460

Gly Lys Pro Gly Gly Lys Pro Glu Gly Gly Gly Glu Gly Lys Pro Gly
465                 470                 475                 480
```

Gly Lys Pro Glu Gly Gly Gly Glu Gly Lys Pro Gly Gly Gly Lys Pro
485 490 495

Gly Glu Gly Gly Lys Pro Glu Gly Gly Gly Lys Pro Gly Glu Gly Gly
500 505 510

Lys Pro Gly Glu Gly Gly Lys Pro Glu Gly Glu Gly Lys Pro Gly Gly
515 520 525

Gly Glu Gly Lys Pro Gly Gly Lys Pro Gly Glu Gly Gly Lys Pro Glu
530 535 540

Gly Gly Gly Glu Gly Lys Pro Gly Gly Lys Pro Gly Gly Glu Gly Gly
545 550 555 560

Gly Lys Pro Glu Gly Gly Lys Pro Gly Glu Gly Gly Lys Pro Glu Gly
565 570 575

<210> 1520
<211> 558
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1520
Gly Glu Gly Lys Pro Gly Gly Lys Pro Gly Ser Gly Gly Gly Lys Pro
1 5 10 15

Gly Glu Gly Gly Lys Pro Gly Ser Gly Glu Gly Lys Pro Gly Gly Lys
20 25 30

Pro Gly Ser Gly Gly Ser Gly Lys Pro Gly Gly Lys Pro Gly Glu Gly
35 40 45

Gly Lys Pro Glu Gly Gly Ser Gly Gly Lys Pro Gly Gly Gly Gly Lys
50 55 60

Pro Gly Gly Lys Pro Gly Gly Glu Gly Ser Gly Lys Pro Gly Gly Lys
65 70 75 80

Pro Glu Gly Gly Gly Lys Pro Glu Gly Gly Ser Gly Gly Lys Pro Gly
85 90 95

Gly Lys Pro Glu Gly Gly Ser Gly Gly Lys Pro Gly Gly Lys Pro Gly
100 105 110

Ser Gly Glu Gly Gly Lys Pro Gly Gly Gly Lys Pro Gly Gly Glu Gly
115 120 125

```
Lys Pro Gly Ser Gly Lys Pro Gly Gly Glu Gly Ser Gly Lys Pro Gly
    130                 135                 140

Gly Lys Pro Glu Gly Gly Ser Gly Gly Lys Pro Gly Gly Lys Pro Glu
145                 150                 155                 160

Gly Gly Ser Gly Gly Lys Pro Gly Gly Ser Gly Lys Pro Gly Gly Lys
                165                 170                 175

Pro Gly Glu Gly Gly Lys Pro Glu Gly Gly Ser Gly Gly Lys Pro Gly
            180                 185                 190

Gly Ser Gly Lys Pro Gly Gly Lys Pro Glu Gly Gly Gly Ser Gly Lys
        195                 200                 205

Pro Gly Gly Lys Pro Gly Glu Gly Gly Lys Pro Gly Ser Gly Glu Gly
    210                 215                 220

Gly Lys Pro Gly Gly Gly Lys Pro Gly Gly Glu Gly Lys Pro Gly Ser
225                 230                 235                 240

Gly Lys Pro Gly Gly Glu Gly Ser Gly Lys Pro Gly Gly Lys Pro Gly
                245                 250                 255

Ser Gly Gly Glu Gly Lys Pro Gly Gly Lys Pro Glu Gly Gly Ser Gly
            260                 265                 270

Gly Lys Pro Gly Gly Gly Lys Pro Gly Gly Glu Gly Lys Pro Gly Ser
        275                 280                 285

Gly Gly Lys Pro Gly Glu Gly Gly Lys Pro Gly Ser Gly Gly Gly Lys
    290                 295                 300

Pro Gly Gly Lys Pro Gly Gly Glu Gly Glu Gly Lys Pro Gly Gly Lys
305                 310                 315                 320

Pro Gly Glu Gly Gly Lys Pro Gly Gly Glu Gly Ser Gly Lys Pro Gly
                325                 330                 335

Gly Gly Gly Lys Pro Gly Gly Lys Pro Gly Gly Glu Gly Gly Lys Pro
            340                 345                 350

Glu Gly Ser Gly Lys Pro Gly Gly Gly Ser Gly Lys Pro Gly Gly Lys
        355                 360                 365

Pro Glu Gly Gly Gly Gly Lys Pro Glu Gly Ser Gly Lys Pro Gly Gly
    370                 375                 380

Gly Gly Lys Pro Glu Gly Ser Gly Lys Pro Gly Gly Gly Lys Pro Glu
385                 390                 395                 400
```

Gly Gly Ser Gly Gly Lys Pro Gly Gly Ser Gly Lys Pro Gly Gly Lys
405 410 415

Pro Gly Glu Gly Gly Gly Lys Pro Glu Gly Ser Gly Lys Pro Gly Gly
420 425 430

Gly Ser Gly Lys Pro Gly Gly Lys Pro Glu Gly Gly Gly Lys Pro Glu
435 440 445

Gly Gly Ser Gly Gly Lys Pro Gly Gly Lys Pro Glu Gly Gly Ser Gly
450 455 460

Gly Lys Pro Gly Gly Lys Pro Gly Gly Glu Gly Ser Gly Lys Pro Gly
465 470 475 480

Gly Lys Pro Gly Ser Gly Glu Gly Gly Lys Pro Gly Gly Lys Pro Gly
485 490 495

Glu Gly Ser Gly Gly Lys Pro Gly Gly Lys Pro Glu Gly Gly Ser Gly
500 505 510

Gly Lys Pro Gly Gly Ser Gly Lys Pro Gly Gly Lys Pro Glu Gly Gly
515 520 525

Gly Ser Gly Lys Pro Gly Gly Lys Pro Gly Glu Gly Gly Lys Pro Gly
530 535 540

Gly Glu Gly Ser Gly Lys Pro Gly Gly Ser Gly Lys Pro Gly
545 550 555

<210> 1521
<211> 577
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1521
Gly Gly Ser Gly Lys Pro Gly Lys Pro Gly Gly Ser Gly Ser Gly Lys
1 5 10 15

Pro Gly Ser Gly Lys Pro Gly Gly Gly Ser Gly Lys Pro Gly Ser Gly
20 25 30

Lys Pro Gly Gly Gly Ser Gly Lys Pro Gly Ser Gly Lys Pro Gly Gly
35 40 45

Gly Ser Gly Lys Pro Gly Ser Gly Lys Pro Gly Gly Gly Gly Lys Pro
50 55 60

Gly Ser Gly Ser Gly Lys Pro Gly Gly Gly Lys Pro Gly Gly Ser Gly
65 70 75 80

```
Gly Lys Pro Gly Gly Gly Ser Gly Lys Pro Gly Lys Pro Gly Ser Gly
                85                  90                  95

Gly Ser Gly Lys Pro Gly Ser Gly Lys Pro Gly Gly Gly Ser Gly Gly
            100                 105                 110

Lys Pro Gly Lys Pro Gly Ser Gly Gly Ser Gly Gly Lys Pro Gly Lys
        115                 120                 125

Pro Gly Ser Gly Gly Gly Ser Gly Lys Pro Gly Lys Pro Gly Ser Gly
    130                 135                 140

Gly Ser Gly Gly Lys Pro Gly Lys Pro Gly Ser Gly Gly Ser Gly Gly
145                 150                 155                 160

Lys Pro Gly Lys Pro Gly Ser Gly Gly Ser Gly Lys Pro Gly Ser Gly
                165                 170                 175

Lys Pro Gly Gly Gly Ser Gly Lys Pro Gly Ser Gly Lys Pro Gly Ser
            180                 185                 190

Gly Gly Ser Gly Lys Pro Gly Lys Pro Gly Ser Gly Gly Ser Gly Lys
        195                 200                 205

Pro Gly Ser Gly Lys Pro Gly Ser Gly Ser Gly Lys Pro Gly Ser Gly
    210                 215                 220

Lys Pro Gly Gly Gly Ser Gly Lys Pro Gly Ser Gly Lys Pro Gly Ser
225                 230                 235                 240

Gly Gly Ser Gly Lys Pro Gly Lys Pro Gly Ser Gly Gly Gly Lys Pro
                245                 250                 255

Gly Ser Gly Ser Gly Lys Pro Gly Gly Gly Lys Pro Gly Ser Gly Ser
            260                 265                 270

Gly Lys Pro Gly Gly Gly Lys Pro Gly Gly Ser Gly Gly Lys Pro Gly
        275                 280                 285

Gly Ser Gly Gly Lys Pro Gly Lys Pro Gly Ser Gly Gly Gly Ser Gly
    290                 295                 300

Lys Pro Gly Lys Pro Gly Ser Gly Gly Gly Ser Gly Lys Pro Gly Lys
305                 310                 315                 320

Pro Gly Gly Ser Gly Ser Gly Lys Pro Gly Ser Gly Lys Pro Gly Gly
                325                 330                 335

Gly Ser Gly Lys Pro Gly Ser Gly Lys Pro Gly Ser Gly Gly Ser Gly
            340                 345                 350
```

Lys Pro Gly Lys Pro Gly Ser Gly Gly Ser Gly Gly Lys Pro Gly Lys
355 360 365

Pro Gly Ser Gly Gly Gly Lys Pro Gly Ser Gly Ser Gly Lys Pro Gly
370 375 380

Gly Gly Lys Pro Gly Ser Gly Ser Gly Lys Pro Gly Gly Gly Lys Pro
385 390 395 400

Gly Ser Gly Ser Gly Lys Pro Gly Gly Gly Lys Pro Gly Ser Gly Ser
405 410 415

Gly Lys Pro Gly Gly Ser Gly Lys Pro Gly Ser Gly Lys Pro Gly Gly
420 425 430

Gly Ser Gly Gly Lys Pro Gly Lys Pro Gly Ser Gly Gly Ser Gly Lys
435 440 445

Pro Gly Ser Gly Lys Pro Gly Ser Gly Gly Ser Gly Lys Pro Gly Lys
450 455 460

Pro Gly Gly Ser Gly Ser Gly Lys Pro Gly Ser Gly Lys Pro Gly Gly
465 470 475 480

Gly Ser Gly Lys Pro Gly Ser Gly Lys Pro Gly Gly Gly Ser Gly Lys
485 490 495

Pro Gly Ser Gly Lys Pro Gly Gly Gly Ser Gly Lys Pro Gly Ser Gly
500 505 510

Lys Pro Gly Gly Gly Gly Lys Pro Gly Ser Gly Ser Gly Lys Pro Gly
515 520 525

Gly Ser Gly Gly Lys Pro Gly Lys Pro Gly Ser Gly Gly Ser Gly Gly
530 535 540

Lys Pro Gly Lys Pro Gly Ser Gly Gly Ser Gly Lys Pro Gly Ser Gly
545 550 555 560

Lys Pro Gly Gly Gly Ser Gly Gly Lys Pro Gly Lys Pro Gly Ser Gly
565 570 575

Gly

<210> 1522
<211> 576
<212> PRT
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1522
Gly Glu Gly Ser Gly Glu Gly Ser Glu Gly Glu Gly Ser Glu Gly Ser
1 5 10 15

Gly Glu Gly Glu Gly Ser Glu Gly Ser Gly Glu Gly Glu Gly Gly Ser
20 25 30

Glu Gly Ser Glu Gly Glu Gly Ser Glu Gly Ser Gly Glu Gly Glu Gly
35 40 45

Gly Glu Gly Ser Gly Glu Gly Glu Gly Ser Gly Glu Gly Ser Glu Gly
50 55 60

Glu Gly Gly Gly Glu Gly Ser Glu Gly Glu Gly Ser Gly Glu Gly Gly
65 70 75 80

Glu Gly Glu Gly Ser Glu Gly Gly Ser Glu Gly Glu Gly Gly Ser Glu
85 90 95

Gly Gly Glu Gly Glu Gly Ser Glu Gly Ser Gly Glu Gly Glu Gly Ser
100 105 110

Glu Gly Gly Ser Glu Gly Glu Gly Ser Glu Gly Gly Ser Glu Gly Glu
115 120 125

Gly Ser Glu Gly Ser Gly Glu Gly Glu Gly Ser Glu Gly Ser Gly Glu
130 135 140

Gly Glu Gly Ser Glu Gly Ser Gly Glu Gly Glu Gly Ser Glu Gly Ser
145 150 155 160

Gly Glu Gly Glu Gly Ser Glu Gly Gly Ser Glu Gly Glu Gly Gly Ser
165 170 175

Glu Gly Ser Glu Gly Glu Gly Ser Gly Glu Gly Ser Glu Gly Glu Gly
180 185 190

Gly Ser Glu Gly Ser Glu Gly Glu Gly Gly Gly Glu Gly Ser Glu Gly
195 200 205

Glu Gly Ser Gly Glu Gly Ser Glu Gly Glu Gly Gly Ser Glu Gly Ser
210 215 220

Glu Gly Glu Gly Gly Ser Glu Gly Ser Glu Gly Glu Gly Gly Glu Gly
225 230 235 240

Ser Gly Glu Gly Glu Gly Ser Glu Gly Ser Gly Glu Gly Glu Gly Ser
245 250 255

```
Gly Glu Gly Ser Glu Gly Glu Gly Ser Glu Gly Ser Gly Glu Gly Glu
            260                 265                 270

Gly Ser Glu Gly Ser Gly Glu Gly Glu Gly Gly Ser Glu Gly Ser Glu
        275                 280                 285

Gly Glu Gly Ser Gly Glu Gly Ser Glu Gly Glu Gly Ser Glu Gly Ser
    290                 295                 300

Gly Glu Gly Glu Gly Ser Glu Gly Ser Gly Glu Gly Glu Gly Gly Ser
305                 310                 315                 320

Glu Gly Ser Glu Gly Glu Gly Gly Ser Glu Gly Ser Glu Gly Glu Gly
                325                 330                 335

Gly Ser Glu Gly Ser Glu Gly Glu Gly Gly Glu Gly Ser Gly Glu Gly
            340                 345                 350

Glu Gly Ser Glu Gly Ser Gly Glu Gly Glu Gly Ser Gly Glu Gly Ser
        355                 360                 365

Glu Gly Glu Gly Ser Glu Gly Ser Gly Glu Gly Glu Gly Ser Glu Gly
    370                 375                 380

Ser Gly Glu Gly Glu Gly Gly Ser Glu Gly Ser Glu Gly Glu Gly Ser
385                 390                 395                 400

Glu Gly Ser Gly Glu Gly Glu Gly Gly Glu Gly Ser Gly Glu Gly Glu
                405                 410                 415

Gly Ser Gly Glu Gly Ser Glu Gly Glu Gly Gly Gly Glu Gly Ser Glu
            420                 425                 430

Gly Glu Gly Ser Glu Gly Ser Gly Glu Gly Glu Gly Ser Glu Gly Ser
        435                 440                 445

Gly Glu Gly Glu Gly Ser Glu Gly Gly Ser Glu Gly Glu Gly Gly Ser
    450                 455                 460

Glu Gly Ser Glu Gly Glu Gly Ser Glu Gly Gly Ser Glu Gly Glu Gly
465                 470                 475                 480

Ser Glu Gly Gly Ser Glu Gly Glu Gly Ser Glu Gly Ser Gly Glu Gly
                485                 490                 495

Glu Gly Ser Glu Gly Ser Gly Glu Gly Glu Gly Ser Gly Glu Gly Ser
            500                 505                 510

Glu Gly Glu Gly Gly Ser Glu Gly Gly Glu Gly Glu Gly Ser Glu Gly
        515                 520                 525
```

Gly Ser Glu Gly Glu Gly Ser Glu Gly Gly Ser Glu Gly Glu Gly Gly
530 535 540

Glu Gly Ser Gly Glu Gly Glu Gly Gly Gly Glu Gly Ser Glu Gly Glu
545 550 555 560

Gly Ser Glu Gly Ser Gly Glu Gly Glu Gly Ser Gly Glu Gly Ser Glu
565 570 575

<210> 1523
<211> 288
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1523
Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
1 5 10 15

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
20 25 30

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
35 40 45

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
50 55 60

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
65 70 75 80

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
85 90 95

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
100 105 110

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
115 120 125

Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
130 135 140

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
145 150 155 160

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
165 170 175

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
180 185 190

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
195 200 205

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
210 215 220

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
225 230 235 240

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
245 250 255

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
260 265 270

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
275 280 285

<210> 1524
<211> 288
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1524
Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
1 5 10 15

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
20 25 30

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
35 40 45

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser
50 55 60

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
65 70 75 80

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
85 90 95

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
100 105 110

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
115 120 125

```
Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
    130                 135                 140

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser
145                 150                 155                 160

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
                165                 170                 175

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
            180                 185                 190

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
        195                 200                 205

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly
    210                 215                 220

Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
225                 230                 235                 240

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
                245                 250                 255

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr
            260                 265                 270

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
        275                 280                 285
```

<210> 1525
<211> 576
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1525

```
Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Gly Gly
1               5                   10                  15

Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Ser Glu Ser Gly Ser Ser
            20                  25                  30

Glu Gly Gly Pro Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro
        35                  40                  45

Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Ser Glu
    50                  55                  60

Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Ser Glu Ser Gly Ser Ser
65                  70                  75                  80
```

```
Glu Gly Gly Pro Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser
                85                  90                  95

Gly Ser Glu Gly Ser Ser Gly Pro Gly Glu Ser Ser Gly Ser Ser Glu
            100                 105                 110

Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Ser Glu Ser Gly Ser Ser
        115                 120                 125

Glu Gly Gly Pro Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro
    130                 135                 140

Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Glu Ser Pro
145                 150                 155                 160

Gly Gly Ser Ser Gly Ser Glu Ser Gly Glu Ser Pro Gly Gly Ser Ser
                165                 170                 175

Gly Ser Glu Ser Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser
            180                 185                 190

Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Gly Gly
        195                 200                 205

Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Gly Gly Glu Pro Ser Glu
    210                 215                 220

Ser Gly Ser Ser Gly Ser Glu Gly Ser Ser Gly Pro Gly Glu Ser Ser
225                 230                 235                 240

Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser Gly Ser Gly Gly
                245                 250                 255

Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Gly Gly Glu Pro Ser Glu
            260                 265                 270

Ser Gly Ser Ser Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser
        275                 280                 285

Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro Gly Glu Ser Pro
    290                 295                 300

Gly Gly Ser Ser Gly Ser Glu Ser Gly Glu Ser Pro Gly Gly Ser Ser
305                 310                 315                 320

Gly Ser Glu Ser Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser
                325                 330                 335

Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser Gly Glu Ser Pro
            340                 345                 350
```

Gly Gly Ser Ser Gly Ser Glu Ser Gly Ser Ser Glu Ser Gly Ser Ser
355 360 365

Glu Gly Gly Pro Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser
370 375 380

Gly Ser Glu Gly Ser Ser Gly Pro Gly Glu Ser Ser Gly Ser Ser Glu
385 390 395 400

Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Gly Gly Glu Pro Ser Glu
405 410 415

Ser Gly Ser Ser Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro
420 425 430

Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Glu Ser Pro
435 440 445

Gly Gly Ser Ser Gly Ser Glu Ser Gly Glu Ser Pro Gly Gly Ser Ser
450 455 460

Gly Ser Glu Ser Gly Ser Ser Glu Ser Gly Ser Ser Glu Gly Gly Pro
465 470 475 480

Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser Gly Ser Ser Glu
485 490 495

Ser Gly Ser Ser Glu Gly Gly Pro Gly Ser Gly Gly Glu Pro Ser Glu
500 505 510

Ser Gly Ser Ser Gly Ser Gly Gly Glu Pro Ser Glu Ser Gly Ser Ser
515 520 525

Gly Glu Ser Pro Gly Gly Ser Ser Gly Ser Glu Ser Gly Ser Glu Gly
530 535 540

Ser Ser Gly Pro Gly Glu Ser Ser Gly Ser Ser Glu Ser Gly Ser Ser
545 550 555 560

Glu Gly Gly Pro Gly Ser Glu Gly Ser Ser Gly Pro Gly Glu Ser Ser
565 570 575

<210> 1526
<211> 577
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1526

```
Ala Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
1               5                   10                  15

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser
            20                  25                  30

Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
        35                  40                  45

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
    50                  55                  60

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
65                  70                  75                  80

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
                85                  90                  95

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro
            100                 105                 110

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr
        115                 120                 125

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
    130                 135                 140

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro
145                 150                 155                 160

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
                165                 170                 175

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            180                 185                 190

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
        195                 200                 205

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
    210                 215                 220

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
225                 230                 235                 240

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
                245                 250                 255

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
            260                 265                 270
```

```
Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
        275                 280                 285

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
    290                 295                 300

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
305                 310                 315                 320

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
                325                 330                 335

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
            340                 345                 350

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
        355                 360                 365

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
    370                 375                 380

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
385                 390                 395                 400

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro
                405                 410                 415

Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            420                 425                 430

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
        435                 440                 445

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
    450                 455                 460

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
465                 470                 475                 480

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
                485                 490                 495

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
            500                 505                 510

Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
        515                 520                 525

Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro
    530                 535                 540
```

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr
545 550 555 560

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
565 570 575

Pro

<210> 1527
<211> 540
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1527
Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser
1 5 10 15

Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser Glu Ser Pro Ser
20 25 30

Gly Thr Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro
35 40 45

Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Thr
50 55 60

Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser Glu Ser Pro Ser
65 70 75 80

Gly Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro
85 90 95

Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser
100 105 110

Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser
115 120 125

Ser Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
130 135 140

Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Pro
145 150 155 160

Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser
165 170 175

Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
180 185 190

```
Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Thr
        195                 200                 205

Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser Glu Ser Pro Ser
    210                 215                 220

Gly Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
225                 230                 235                 240

Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser
                245                 250                 255

Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Thr Pro Glu Ser Gly
            260                 265                 270

Ser Ala Ser Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
        275                 280                 285

Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Thr
    290                 295                 300

Pro Glu Ser Gly Ser Ala Ser Pro Gly Thr Ser Thr Pro Glu Ser Gly
305                 310                 315                 320

Ser Ala Ser Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
                325                 330                 335

Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser
            340                 345                 350

Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu
        355                 360                 365

Ser Pro Gly Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
    370                 375                 380

Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser
385                 390                 395                 400

Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser
                405                 410                 415

Gly Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
            420                 425                 430

Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser
        435                 440                 445

Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly
    450                 455                 460
```

Ser Ala Ser Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro
465 470 475 480

Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro
485 490 495

Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu
500 505 510

Ser Pro Gly Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
515 520 525

Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
530 535 540

<210> 1528
<211> 504
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1528
Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
1 5 10 15

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
20 25 30

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
35 40 45

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Asn Pro
50 55 60

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
65 70 75 80

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
85 90 95

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
100 105 110

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
115 120 125

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
130 135 140

```
Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
145                 150                 155                 160

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
                165                 170                 175

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
            180                 185                 190

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Asn Pro
        195                 200                 205

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
    210                 215                 220

Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
225                 230                 235                 240

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro
                245                 250                 255

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
            260                 265                 270

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
        275                 280                 285

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
    290                 295                 300

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
305                 310                 315                 320

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                325                 330                 335

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
            340                 345                 350

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
        355                 360                 365

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    370                 375                 380

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
385                 390                 395                 400

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
                405                 410                 415
```

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
420 425 430

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
435 440 445

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
450 455 460

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
465 470 475 480

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
485 490 495

Gly Thr Ser Ser Thr Gly Ser Pro
500

<210> 1529
<211> 816
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1529
Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
35 40 45

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
50 55 60

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
65 70 75 80

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
85 90 95

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
100 105 110

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
115 120 125

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
130 135 140

```
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
145                 150                 155                 160

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
                165                 170                 175

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            180                 185                 190

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
        195                 200                 205

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
    210                 215                 220

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
225                 230                 235                 240

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
                245                 250                 255

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
            260                 265                 270

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        275                 280                 285

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
    290                 295                 300

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
305                 310                 315                 320

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
                325                 330                 335

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
            340                 345                 350

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
        355                 360                 365

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    370                 375                 380

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
385                 390                 395                 400

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
                405                 410                 415
```

```
Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            420                 425                 430

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
        435                 440                 445

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
    450                 455                 460

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
465                 470                 475                 480

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
                485                 490                 495

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
            500                 505                 510

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
        515                 520                 525

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala
    530                 535                 540

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
545                 550                 555                 560

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
                565                 570                 575

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
            580                 585                 590

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
        595                 600                 605

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
    610                 615                 620

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
625                 630                 635                 640

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
                645                 650                 655

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
            660                 665                 670

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
        675                 680                 685
```

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
690 695 700

Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
705 710 715 720

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
725 730 735

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
740 745 750

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
755 760 765

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
770 775 780

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
785 790 795 800

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
805 810 815

<210> 1530
<211> 875
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<220>
<221> MOD_RES
<222> (430)..(432)
<223> Any amino acid

<220>
<221> MOD_RES
<222> (443)..(446)
<223> Any amino acid

<400> 1530
Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Pro
1 5 10 15

Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser
20 25 30

Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
35 40 45

Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Thr Ser Thr

```
    50                  55                  60

Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser Glu Ser Pro Ser
65                  70                  75                  80

Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
                85                  90                  95

Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser
            100                 105                 110

Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser
        115                 120                 125

Ser Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro
    130                 135                 140

Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro
145                 150                 155                 160

Ser Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser
                165                 170                 175

Ser Thr Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro
            180                 185                 190

Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Thr Ser Thr
        195                 200                 205

Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser Glu Ser Pro Ser
    210                 215                 220

Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
225                 230                 235                 240

Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser
                245                 250                 255

Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Thr Pro Glu Ser Gly
            260                 265                 270

Ser Ala Ser Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
        275                 280                 285

Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser
    290                 295                 300

Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro Ser Gly Glu Ser
305                 310                 315                 320

Ser Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
```

325 330 335

Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser
340 345 350

Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser Ser Thr Ala Glu
355 360 365

Ser Pro Gly Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro
370 375 380

Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser
385 390 395 400

Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser
405 410 415

Gly Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Pro Xaa Xaa Xaa
420 425 430

Gly Ala Ser Ala Ser Gly Ala Pro Ser Thr Xaa Xaa Xaa Xaa Ser Glu
435 440 445

Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly
450 455 460

Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly
465 470 475 480

Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu
485 490 495

Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly
500 505 510

Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly
515 520 525

Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Pro Ser
530 535 540

Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser
545 550 555 560

Pro Gly Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly
565 570 575

Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser Glu
580 585 590

Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly

595 600 605

Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly
610 615 620

Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Thr Pro
625 630 635 640

Glu Ser Gly Ser Ala Ser Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser
645 650 655

Ala Ser Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly
660 665 670

Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser Ser
675 680 685

Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly
690 695 700

Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly
705 710 715 720

Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser Ser
725 730 735

Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser
740 745 750

Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly
755 760 765

Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Pro Ser
770 775 780

Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser
785 790 795 800

Thr Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly
805 810 815

Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro Ser
820 825 830

Gly Glu Ser Ser Thr Ala Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
835 840 845

Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
850 855 860

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro

865 870 875

<210> 1531
<211> 864
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1531
Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
1 5 10 15

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
20 25 30

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
35 40 45

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
50 55 60

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
65 70 75 80

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
85 90 95

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
100 105 110

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
115 120 125

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
130 135 140

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
145 150 155 160

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
165 170 175

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
180 185 190

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
195 200 205

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
210 215 220

```
Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
225                 230                 235                 240

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro
                245                 250                 255

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
            260                 265                 270

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
        275                 280                 285

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
    290                 295                 300

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
305                 310                 315                 320

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                325                 330                 335

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
            340                 345                 350

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
        355                 360                 365

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    370                 375                 380

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
385                 390                 395                 400

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
                405                 410                 415

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
            420                 425                 430

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
        435                 440                 445

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
    450                 455                 460

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
465                 470                 475                 480

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
                485                 490                 495
```

```
Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
            500                 505                 510

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
        515                 520                 525

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
    530                 535                 540

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
545                 550                 555                 560

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                565                 570                 575

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
            580                 585                 590

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
        595                 600                 605

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
    610                 615                 620

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
625                 630                 635                 640

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
                645                 650                 655

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            660                 665                 670

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
        675                 680                 685

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
    690                 695                 700

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
705                 710                 715                 720

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
                725                 730                 735

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
            740                 745                 750

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
        755                 760                 765
```

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
770 775 780

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
785 790 795 800

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
805 810 815

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
820 825 830

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
835 840 845

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
850 855 860

<210> 1532
<211> 868
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1532
Gly Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
1 5 10 15

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
20 25 30

Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala
35 40 45

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
50 55 60

Gly Ser Asn Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
65 70 75 80

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
85 90 95

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
100 105 110

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
115 120 125

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser

130 135 140

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
145 150 155 160

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro
165 170 175

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
180 185 190

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
195 200 205

Gly Ser Asn Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
210 215 220

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala
225 230 235 240

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
245 250 255

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
260 265 270

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
275 280 285

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
290 295 300

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
305 310 315 320

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
325 330 335

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
340 345 350

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
355 360 365

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
370 375 380

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
385 390 395 400

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr

405 410 415

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
420 425 430

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
435 440 445

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
450 455 460

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
465 470 475 480

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
485 490 495

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
500 505 510

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
515 520 525

Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
530 535 540

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
545 550 555 560

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
565 570 575

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
580 585 590

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
595 600 605

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
610 615 620

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
625 630 635 640

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
645 650 655

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
660 665 670

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro

675 680 685

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
690 695 700

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
705 710 715 720

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
725 730 735

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
740 745 750

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
755 760 765

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
770 775 780

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
785 790 795 800

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
805 810 815

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
820 825 830

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
835 840 845

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
850 855 860

Thr Gly Ser Pro
865

<210> 1533
<211> 875
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1533
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
1 5 10 15

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
20 25 30

```
Ser Thr Glu Glu Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro
        35                  40                  45

Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser
    50                  55                  60

Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser
65                  70                  75                  80

Gly Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
                85                  90                  95

Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Glu Pro
            100                 105                 110

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
        115                 120                 125

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
    130                 135                 140

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
145                 150                 155                 160

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
                165                 170                 175

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            180                 185                 190

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
        195                 200                 205

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
    210                 215                 220

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
225                 230                 235                 240

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
                245                 250                 255

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
            260                 265                 270

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        275                 280                 285

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
    290                 295                 300
```

```
Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
305                 310                 315                 320

Ser Thr Glu Glu Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
                325                 330                 335

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
            340                 345                 350

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Ser Thr Glu Pro Ser Glu
        355                 360                 365

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    370                 375                 380

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
385                 390                 395                 400

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
                405                 410                 415

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            420                 425                 430

Gly Ala Ser Ala Ser Gly Ala Pro Ser Thr Gly Gly Thr Ser Glu Ser
        435                 440                 445

Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
    450                 455                 460

Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
465                 470                 475                 480

Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser Glu
                485                 490                 495

Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser
            500                 505                 510

Thr Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
        515                 520                 525

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser
    530                 535                 540

Ala Ser Thr Gly Thr Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
545                 550                 555                 560

Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
                565                 570                 575
```

```
Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Thr Ser Ser
            580                 585                 590

Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser
        595                 600                 605

Pro Gly Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly
    610                 615                 620

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala
625                 630                 635                 640

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
                645                 650                 655

Ser Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly
            660                 665                 670

Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser Glu
        675                 680                 685

Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    690                 695                 700

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
705                 710                 715                 720

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Ser Thr Pro
                725                 730                 735

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
            740                 745                 750

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
        755                 760                 765

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
    770                 775                 780

Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
785                 790                 795                 800

Thr Glu Glu Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
                805                 810                 815

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly
            820                 825                 830

Thr Ser Ser Thr Gly Ser Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
        835                 840                 845
```

Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
850 855 860

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
865 870 875

<210> 1534
<211> 913
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1534
Met Ala Glu Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Pro
1 5 10 15

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
20 25 30

Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
35 40 45

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
50 55 60

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser
65 70 75 80

Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
85 90 95

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
100 105 110

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
115 120 125

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
130 135 140

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro
145 150 155 160

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr
165 170 175

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
180 185 190

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro

```
        195                 200                 205

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
    210                 215                 220

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
225                 230                 235                 240

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
                245                 250                 255

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
            260                 265                 270

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
        275                 280                 285

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
    290                 295                 300

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
305                 310                 315                 320

Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
                325                 330                 335

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
            340                 345                 350

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
        355                 360                 365

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
    370                 375                 380

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
385                 390                 395                 400

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
                405                 410                 415

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            420                 425                 430

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
        435                 440                 445

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro
    450                 455                 460

Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
```

```
465                 470                 475                 480

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
                485                 490                 495

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
            500                 505                 510

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
        515                 520                 525

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
    530                 535                 540

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
545                 550                 555                 560

Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
                565                 570                 575

Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro
            580                 585                 590

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr
        595                 600                 605

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
    610                 615                 620

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
625                 630                 635                 640

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
                645                 650                 655

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
            660                 665                 670

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
        675                 680                 685

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro
    690                 695                 700

Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
705                 710                 715                 720

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser
                725                 730                 735

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro
```

740 745 750

Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
755 760 765

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
770 775 780

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
785 790 795 800

Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
805 810 815

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu
820 825 830

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro
835 840 845

Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
850 855 860

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu
865 870 875 880

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
885 890 895

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
900 905 910

Pro

<210> 1535
<211> 924
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1535
Met Ala Glu Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ala Ser
1 5 10 15

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly
20 25 30

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
35 40 45

```
Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu
    50                  55                  60

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro
65                  70                  75                  80

Thr Ser Thr Glu Glu Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly
                85                  90                  95

Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr
            100                 105                 110

Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro
        115                 120                 125

Ser Gly Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser
    130                 135                 140

Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Glu
145                 150                 155                 160

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
                165                 170                 175

Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
            180                 185                 190

Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
        195                 200                 205

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser
    210                 215                 220

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
225                 230                 235                 240

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
                245                 250                 255

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
            260                 265                 270

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
        275                 280                 285

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
    290                 295                 300

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
305                 310                 315                 320
```

```
Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
                325                 330                 335

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu
            340                 345                 350

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro
        355                 360                 365

Thr Ser Thr Glu Glu Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
    370                 375                 380

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser
385                 390                 395                 400

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Ser Thr Glu Pro Ser
                405                 410                 415

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
            420                 425                 430

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro
        435                 440                 445

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro
    450                 455                 460

Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
465                 470                 475                 480

Pro Gly Ala Ser Ala Ser Gly Ala Pro Ser Thr Gly Gly Thr Ser Glu
                485                 490                 495

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
            500                 505                 510

Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
        515                 520                 525

Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser
    530                 535                 540

Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser
545                 550                 555                 560

Ser Thr Ala Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
                565                 570                 575

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
            580                 585                 590
```

```
Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
        595                 600                 605

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
    610                 615                 620

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Thr Ser
625                 630                 635                 640

Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser Ser Thr Ala Glu
                645                 650                 655

Ser Pro Gly Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro
            660                 665                 670

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
        675                 680                 685

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu
    690                 695                 700

Gly Ser Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro
705                 710                 715                 720

Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser
                725                 730                 735

Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
            740                 745                 750

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
        755                 760                 765

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Ser Thr
    770                 775                 780

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
785                 790                 795                 800

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                805                 810                 815

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
            820                 825                 830

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
        835                 840                 845

Ser Thr Glu Glu Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
    850                 855                 860
```

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
865 870 875 880

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Ser Glu Ser Ala Thr Pro
885 890 895

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
900 905 910

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
915 920

<210> 1536
<211> 1318
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1536
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
1 5 10 15

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
20 25 30

Ser Thr Glu Glu Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro
35 40 45

Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser
50 55 60

Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser
65 70 75 80

Gly Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
85 90 95

Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Glu Pro
100 105 110

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
115 120 125

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
130 135 140

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
145 150 155 160

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu

```
                165                 170                 175

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            180                 185                 190

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
        195                 200                 205

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
    210                 215                 220

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
225                 230                 235                 240

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
                245                 250                 255

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
            260                 265                 270

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        275                 280                 285

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
    290                 295                 300

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
305                 310                 315                 320

Ser Thr Glu Glu Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
                325                 330                 335

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
            340                 345                 350

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Ser Thr Glu Pro Ser Glu
        355                 360                 365

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    370                 375                 380

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
385                 390                 395                 400

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
                405                 410                 415

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            420                 425                 430

Gly Pro Glu Pro Thr Gly Pro Ala Pro Ser Gly Gly Ser Glu Pro Ala
```

435 440 445

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
450 455 460

Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
465 470 475 480

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly
485 490 495

Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser
500 505 510

Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
515 520 525

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
530 535 540

Ser Pro Thr Ser Thr Glu Glu Gly Ser Thr Ser Ser Thr Ala Glu Ser
545 550 555 560

Pro Gly Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly
565 570 575

Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser Glu
580 585 590

Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly
595 600 605

Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly
610 615 620

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
625 630 635 640

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
645 650 655

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
660 665 670

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser
675 680 685

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
690 695 700

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly

705 710 715 720

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Pro Ser
725 730 735

Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser
740 745 750

Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly
755 760 765

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
770 775 780

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
785 790 795 800

Ser Ala Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
805 810 815

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
820 825 830

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
835 840 845

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
850 855 860

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Ala Ser
865 870 875 880

Gly Ala Pro Ser Thr Gly Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr
885 890 895

Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr
900 905 910

Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Glu Ser Ala
915 920 925

Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
930 935 940

Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
945 950 955 960

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser
965 970 975

Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly

980 985 990

Ser Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Thr
995 1000 1005

Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Pro Ser
1010 1015 1020

Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
1025 1030 1035

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1040 1045 1050

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
1055 1060 1065

Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu
1070 1075 1080

Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly
1085 1090 1095

Ser Ala Ser Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
1100 1105 1110

Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
1115 1120 1125

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
1130 1135 1140

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
1145 1150 1155

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1160 1165 1170

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala
1175 1180 1185

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro
1190 1195 1200

Ser Gly Ala Thr Gly Ser Pro Gly Ser Thr Ser Glu Ser Pro Ser
1205 1210 1215

Gly Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala
1220 1225 1230

Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Ser

1235 1240 1245

Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly
1250 1255 1260

Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
1265 1270 1275

Ser Ser Ser Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu
1280 1285 1290

Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr
1295 1300 1305

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
1310 1315

<210> 1537
<211> 1874
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1537
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Gly
115 120 125

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
130 135 140

```
Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
145                 150                 155                 160

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
                165                 170                 175

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
            180                 185                 190

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
        195                 200                 205

Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
    210                 215                 220

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
225                 230                 235                 240

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
                245                 250                 255

Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            260                 265                 270

Gly Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys
        275                 280                 285

Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr
    290                 295                 300

Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly
305                 310                 315                 320

Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln
                325                 330                 335

Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys
            340                 345                 350

Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala
        355                 360                 365

Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val
    370                 375                 380

Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr
385                 390                 395                 400

Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe
                405                 410                 415
```

```
Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu
            420                 425                 430

Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp
        435                 440                 445

Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp
    450                 455                 460

Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln
465                 470                 475                 480

Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu
                485                 490                 495

Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro
            500                 505                 510

Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp
        515                 520                 525

Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu
    530                 535                 540

Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn
545                 550                 555                 560

Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala
                565                 570                 575

Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser
            580                 585                 590

Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu
        595                 600                 605

Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His
    610                 615                 620

Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly
625                 630                 635                 640

Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys
                645                 650                 655

Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro
            660                 665                 670

Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp
        675                 680                 685
```

```
Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser
    690                 695                 700

Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile
705                 710                 715                 720

Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn
                725                 730                 735

Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro
            740                 745                 750

Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe
        755                 760                 765

Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr
    770                 775                 780

Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser
785                 790                 795                 800

Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu
                805                 810                 815

Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly
            820                 825                 830

Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met
        835                 840                 845

Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr
    850                 855                 860

Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn
865                 870                 875                 880

Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Gly Thr Ser Glu
                885                 890                 895

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
            900                 905                 910

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        915                 920                 925

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
    930                 935                 940

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
945                 950                 955                 960
```

```
Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
               965                 970                 975

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
           980                 985                 990

Ala Thr Ser Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
       995                1000                1005

Glu Ser  Gly Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    1010                1015                1020

Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr
    1025                1030                1035

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Thr  Ser Glu Ser
    1040                1045                1050

Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1055                1060                1065

Glu Ser  Gly Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1070                1075                1080

Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Ser
    1085                1090                1095

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Ser  Pro Ala Gly
    1100                1105                1110

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    1115                1120                1125

Gly Ser  Ala Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    1130                1135                1140

Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr
    1145                1150                1155

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Thr  Ser Thr Glu
    1160                1165                1170

Pro Ser  Glu Gly Ser Ala Pro  Gly Pro Pro Val Leu  Lys Arg His
    1175                1180                1185

Gln Arg  Glu Ile Thr Arg Thr  Thr Leu Gln Ser Asp  Gln Glu Glu
    1190                1195                1200

Ile Asp  Tyr Asp Asp Thr Ile  Ser Val Glu Met Lys  Lys Glu Asp
    1205                1210                1215
```

```
Phe Asp  Ile Tyr Asp Glu Asp  Glu Asn Gln Ser Pro  Arg Ser Phe
    1220                 1225                 1230

Gln Lys  Lys Thr Arg His Tyr  Phe Ile Ala Ala Val  Glu Arg Leu
    1235                 1240                 1245

Trp Asp  Tyr Gly Met Ser Ser  Ser Pro His Val Leu  Arg Asn Arg
    1250                 1255                 1260

Ala Gln  Ser Gly Ser Val Pro  Gln Phe Lys Lys Val  Val Phe Gln
    1265                 1270                 1275

Glu Phe  Thr Asp Gly Ser Phe  Thr Gln Pro Leu Tyr  Arg Gly Glu
    1280                 1285                 1290

Leu Asn  Glu His Leu Gly Leu  Leu Gly Pro Tyr Ile  Arg Ala Glu
    1295                 1300                 1305

Val Glu  Asp Asn Ile Met Val  Thr Phe Arg Asn Gln  Ala Ser Arg
    1310                 1315                 1320

Pro Tyr  Ser Phe Tyr Ser Ser  Leu Ile Ser Tyr Glu  Glu Asp Gln
    1325                 1330                 1335

Arg Gln  Gly Ala Glu Pro Arg  Lys Asn Phe Val Lys  Pro Asn Glu
    1340                 1345                 1350

Thr Lys  Thr Tyr Phe Trp Lys  Val Gln His His Met  Ala Pro Thr
    1355                 1360                 1365

Lys Asp  Glu Phe Asp Cys Lys  Ala Trp Ala Tyr Phe  Ser Asp Val
    1370                 1375                 1380

Asp Leu  Glu Lys Asp Val His  Ser Gly Leu Ile Gly  Pro Leu Leu
    1385                 1390                 1395

Val Cys  His Thr Asn Thr Leu  Asn Pro Ala His Gly  Arg Gln Val
    1400                 1405                 1410

Thr Val  Gln Glu Phe Ala Leu  Phe Phe Thr Ile Phe  Asp Glu Thr
    1415                 1420                 1425

Lys Ser  Trp Tyr Phe Thr Glu  Asn Met Glu Arg Asn  Cys Arg Ala
    1430                 1435                 1440

Pro Cys  Asn Ile Gln Met Glu  Asp Pro Thr Phe Lys  Glu Asn Tyr
    1445                 1450                 1455

Arg Phe  His Ala Ile Asn Gly  Tyr Ile Met Asp Thr  Leu Pro Gly
    1460                 1465                 1470
```

```
Leu Val  Met Ala Gln Asp Gln  Arg Ile Arg Trp Tyr  Leu Leu Ser
    1475                 1480                 1485

Met Gly  Ser Asn Glu Asn Ile  His Ser Ile His Phe  Ser Gly His
    1490                 1495                 1500

Val Phe  Thr Val Arg Lys Lys  Glu Glu Tyr Lys Met  Ala Leu Tyr
    1505                 1510                 1515

Asn Leu  Tyr Pro Gly Val Phe  Glu Thr Val Glu Met  Leu Pro Ser
    1520                 1525                 1530

Lys Ala  Gly Ile Trp Arg Val  Glu Cys Leu Ile Gly  Glu His Leu
    1535                 1540                 1545

His Ala  Gly Met Ser Thr Leu  Phe Leu Val Tyr Ser  Asn Lys Cys
    1550                 1555                 1560

Gln Thr  Pro Leu Gly Met Ala  Ser Gly His Ile Arg  Asp Phe Gln
    1565                 1570                 1575

Ile Thr  Ala Ser Gly Gln Tyr  Gly Gln Trp Ala Pro  Lys Leu Ala
    1580                 1585                 1590

Arg Leu  His Tyr Ser Gly Ser  Ile Asn Ala Trp Ser  Thr Lys Glu
    1595                 1600                 1605

Pro Phe  Ser Trp Ile Lys Val  Asp Leu Leu Ala Pro  Met Ile Ile
    1610                 1615                 1620

His Gly  Ile Lys Thr Gln Gly  Ala Arg Gln Lys Phe  Ser Ser Leu
    1625                 1630                 1635

Tyr Ile  Ser Gln Phe Ile Ile  Met Tyr Ser Leu Asp  Gly Lys Lys
    1640                 1645                 1650

Trp Gln  Thr Tyr Arg Gly Asn  Ser Thr Gly Thr Leu  Met Val Phe
    1655                 1660                 1665

Phe Gly  Asn Val Asp Ser Ser  Gly Ile Lys His Asn  Ile Phe Asn
    1670                 1675                 1680

Pro Pro  Ile Ile Ala Arg Tyr  Ile Arg Leu His Pro  Thr His Tyr
    1685                 1690                 1695

Ser Ile  Arg Ser Thr Leu Arg  Met Glu Leu Met Gly  Cys Asp Leu
    1700                 1705                 1710

Asn Ser  Cys Ser Met Pro Leu  Gly Met Glu Ser Lys  Ala Ile Ser
    1715                 1720                 1725
```

```
Asp Ala  Gln Ile Thr Ala Ser  Ser Tyr Phe Thr Asn  Met Phe Ala
    1730                1735                1740

Thr Trp  Ser Pro Ser Lys Ala  Arg Leu His Leu Gln  Gly Arg Ser
    1745                1750                1755

Asn Ala  Trp Arg Pro Gln Val  Asn Asn Pro Lys Glu  Trp Leu Gln
    1760                1765                1770

Val Asp  Phe Gln Lys Thr Met  Lys Val Thr Gly Val  Thr Thr Gln
    1775                1780                1785

Gly Val  Lys Ser Leu Leu Thr  Ser Met Tyr Val Lys  Glu Phe Leu
    1790                1795                1800

Ile Ser  Ser Ser Gln Asp Gly  His Gln Trp Thr Leu  Phe Phe Gln
    1805                1810                1815

Asn Gly  Lys Val Lys Val Phe  Gln Gly Asn Gln Asp  Ser Phe Thr
    1820                1825                1830

Pro Val  Val Asn Ser Leu Asp  Pro Pro Leu Leu Thr  Arg Tyr Leu
    1835                1840                1845

Arg Ile  His Pro Gln Ser Trp  Val His Gln Ile Ala  Leu Arg Met
    1850                1855                1860

Glu Val  Leu Gly Cys Glu Ala  Gln Asp Leu Tyr
    1865                1870
```

<210> 1538
<211> 2162
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1538

```
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1               5                  10                  15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
```

65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp

340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Gly Gly Ser Pro Ala Gly Ser Pro Thr
370 375 380

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
385 390 395 400

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
405 410 415

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
420 425 430

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
435 440 445

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
450 455 460

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
465 470 475 480

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
485 490 495

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
500 505 510

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
515 520 525

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
530 535 540

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
545 550 555 560

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
565 570 575

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
580 585 590

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
595 600 605

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu

610 615 620

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
625 630 635 640

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu
645 650 655

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
660 665 670

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
675 680 685

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
690 695 700

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
705 710 715 720

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
725 730 735

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
740 745 750

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
755 760 765

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
770 775 780

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
785 790 795 800

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
805 810 815

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
820 825 830

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
835 840 845

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
850 855 860

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
865 870 875 880

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala

885 890 895

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr
900 905 910

Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
915 920 925

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
930 935 940

Glu Pro Ser Glu Gly Ser Ala Pro Gly Lys Lys His Pro Lys Thr Trp
945 950 955 960

Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu
965 970 975

Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn
980 985 990

Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala
995 1000 1005

Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu
1010 1015 1020

Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr
1025 1030 1035

Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile
1040 1045 1050

Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
1055 1060 1065

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro
1070 1075 1080

Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly
1085 1090 1095

Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser
1100 1105 1110

Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro
1115 1120 1125

Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln
1130 1135 1140

Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp

1145 1150 1155

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu
1160 1165 1170

Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala
1175 1180 1185

Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu
1190 1195 1200

Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
1205 1210 1215

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly
1220 1225 1230

Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu
1235 1240 1245

Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro
1250 1255 1260

Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg
1265 1270 1275

Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr
1280 1285 1290

Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
1295 1300 1305

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn
1310 1315 1320

Gly Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr
1325 1330 1335

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1340 1345 1350

Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
1355 1360 1365

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
1370 1375 1380

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser
1385 1390 1395

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala

1400 1405 1410

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu
1415 1420 1425

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1430 1435 1440

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr
1445 1450 1455

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Pro Pro Val Leu
1460 1465 1470

Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp
1475 1480 1485

Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys
1490 1495 1500

Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
1505 1510 1515

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val
1520 1525 1530

Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu
1535 1540 1545

Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val
1550 1555 1560

Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
1565 1570 1575

Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile
1580 1585 1590

Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln
1595 1600 1605

Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu
1610 1615 1620

Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys
1625 1630 1635

Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His Met
1640 1645 1650

Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe

1655 1660 1665

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly
1670 1675 1680

Pro Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly
1685 1690 1695

Arg Gln Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe
1700 1705 1710

Asp Glu Thr Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn
1715 1720 1725

Cys Arg Ala Pro Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys
1730 1735 1740

Glu Asn Tyr Arg Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr
1745 1750 1755

Leu Pro Gly Leu Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr
1760 1765 1770

Leu Leu Ser Met Gly Ser Asn Glu Asn Ile His Ser Ile His Phe
1775 1780 1785

Ser Gly His Val Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met
1790 1795 1800

Ala Leu Tyr Asn Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met
1805 1810 1815

Leu Pro Ser Lys Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly
1820 1825 1830

Glu His Leu His Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser
1835 1840 1845

Asn Lys Cys Gln Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg
1850 1855 1860

Asp Phe Gln Ile Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro
1865 1870 1875

Lys Leu Ala Arg Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser
1880 1885 1890

Thr Lys Glu Pro Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro
1895 1900 1905

Met Ile Ile His Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe

1910 1915 1920

Ser Ser Leu Tyr Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp
1925 1930 1935

Gly Lys Lys Trp Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu
1940 1945 1950

Met Val Phe Phe Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn
1955 1960 1965

Ile Phe Asn Pro Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro
1970 1975 1980

Thr His Tyr Ser Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly
1985 1990 1995

Cys Asp Leu Asn Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys
2000 2005 2010

Ala Ile Ser Asp Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn
2015 2020 2025

Met Phe Ala Thr Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln
2030 2035 2040

Gly Arg Ser Asn Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu
2045 2050 2055

Trp Leu Gln Val Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val
2060 2065 2070

Thr Thr Gln Gly Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys
2075 2080 2085

Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu
2090 2095 2100

Phe Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp
2105 2110 2115

Ser Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr
2120 2125 2130

Arg Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile Ala
2135 2140 2145

Leu Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr
2150 2155 2160

<210> 1539

<211> 2449
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1539
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

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Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510
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Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780
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Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Gly Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly
            900                 905                 910

Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Pro Ser
        915                 920                 925

Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser
    930                 935                 940

Pro Gly Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly
945                 950                 955                 960

Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro Ser
                965                 970                 975

Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser
            980                 985                 990

Ala Ser Pro Gly Ser Thr Ser Ser  Thr Ala Glu Ser Pro  Gly Pro Gly
        995                 1000                1005

Thr Ser  Pro Ser Gly Glu Ser  Ser Thr Ala Pro Gly  Thr Ser Pro
    1010                1015                1020

Ser Gly  Glu Ser Ser Thr Ala  Pro Gly Thr Ser Pro  Ser Gly Glu
    1025                1030                1035

Ser Ser  Thr Ala Pro Gly Glu  Glu Asp Gln Arg Gln  Gly Ala Glu
    1040                1045                1050
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Pro Arg  Lys Asn Phe Val Lys  Pro Asn Glu Thr Lys  Thr Tyr Phe
    1055                1060                1065

Trp Lys  Val Gln His His Met  Ala Pro Thr Lys Asp  Glu Phe Asp
    1070                1075                1080

Cys Lys  Ala Trp Ala Tyr Phe  Ser Asp Val Asp Leu  Glu Lys Asp
    1085                1090                1095

Val His  Ser Gly Leu Ile Gly  Pro Leu Leu Val Cys  His Thr Asn
    1100                1105                1110

Thr Leu  Asn Pro Ala His Gly  Arg Gln Val Thr Val  Gln Glu Phe
    1115                1120                1125

Ala Leu  Phe Phe Thr Ile Phe  Asp Glu Thr Lys Ser  Trp Tyr Phe
    1130                1135                1140

Thr Glu  Asn Met Glu Arg Asn  Cys Arg Ala Pro Cys  Asn Ile Gln
    1145                1150                1155

Met Glu  Asp Pro Thr Phe Lys  Glu Asn Tyr Arg Phe  His Ala Ile
    1160                1165                1170

Asn Gly  Tyr Ile Met Asp Thr  Leu Pro Gly Leu Val  Met Ala Gln
    1175                1180                1185

Asp Gln  Arg Ile Arg Trp Tyr  Leu Leu Ser Met Gly  Ser Asn Glu
    1190                1195                1200

Asn Ile  His Ser Ile His Phe  Ser Gly His Val Phe  Thr Val Arg
    1205                1210                1215

Lys Lys  Glu Glu Tyr Lys Met  Ala Leu Tyr Asn Leu  Tyr Pro Gly
    1220                1225                1230

Val Phe  Glu Thr Val Glu Met  Leu Pro Ser Lys Ala  Gly Ile Trp
    1235                1240                1245

Arg Val  Glu Cys Leu Ile Gly  Glu His Leu His Ala  Gly Met Ser
    1250                1255                1260

Thr Leu  Phe Leu Val Tyr Ser  Asn Lys Cys Gln Thr  Pro Leu Gly
    1265                1270                1275

Met Ala  Ser Gly His Ile Arg  Asp Phe Gln Ile Thr  Ala Ser Gly
    1280                1285                1290

Gln Tyr  Gly Gln Trp Ala Pro  Lys Leu Ala Arg Leu  His Tyr Ser
    1295                1300                1305
```

```
Gly Ser  Ile Asn Ala Trp Ser  Thr Lys Glu Pro Phe  Ser Trp Ile
    1310                1315                1320

Lys Val  Asp Leu Leu Ala Pro  Met Ile Ile His Gly  Ile Lys Thr
    1325                1330                1335

Gln Gly  Ala Arg Gln Lys Phe  Ser Ser Leu Tyr Ile  Ser Gln Phe
    1340                1345                1350

Ile Ile  Met Tyr Ser Leu Asp  Gly Lys Lys Trp Gln  Thr Tyr Arg
    1355                1360                1365

Gly Asn  Ser Thr Gly Thr Leu  Met Val Phe Phe Gly  Asn Val Asp
    1370                1375                1380

Ser Ser  Gly Ile Lys His Asn  Ile Phe Asn Pro Pro  Ile Ile Ala
    1385                1390                1395

Arg Tyr  Ile Arg Leu His Pro  Thr His Tyr Ser Ile  Arg Ser Thr
    1400                1405                1410

Leu Arg  Met Glu Leu Met Gly  Cys Asp Leu Asn Ser  Cys Ser Met
    1415                1420                1425

Pro Leu  Gly Met Glu Ser Lys  Ala Ile Ser Asp Ala  Gln Ile Thr
    1430                1435                1440

Ala Ser  Ser Tyr Phe Thr Asn  Met Phe Ala Thr Trp  Ser Pro Ser
    1445                1450                1455

Lys Ala  Arg Leu His Leu Gln  Gly Arg Ser Asn Ala  Trp Arg Pro
    1460                1465                1470

Gln Val  Asn Asn Pro Lys Glu  Trp Leu Gln Val Asp  Phe Gln Lys
    1475                1480                1485

Thr Met  Lys Val Thr Gly Val  Thr Thr Gln Gly Val  Lys Ser Leu
    1490                1495                1500

Leu Thr  Ser Met Tyr Val Lys  Glu Phe Leu Ile Ser  Ser Ser Gln
    1505                1510                1515

Asp Gly  His Gln Trp Thr Leu  Phe Phe Gln Asn Gly  Lys Val Lys
    1520                1525                1530

Val Phe  Gln Gly Asn Gln Asp  Ser Phe Thr Pro Val  Val Asn Ser
    1535                1540                1545

Leu Asp  Pro Pro Leu Leu Thr  Arg Tyr Leu Arg Ile  His Pro Gln
    1550                1555                1560
```

```
Ser Trp  Val His Gln Ile Ala  Leu Arg Met Glu Val  Leu Gly Cys
    1565                1570                1575

Glu Ala  Gln Asp Leu Tyr Gly  Gly Ser Pro Ala Gly  Ser Pro Thr
    1580                1585                1590

Ser Thr  Glu Glu Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1595                1600                1605

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    1610                1615                1620

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Thr Glu
    1625                1630                1635

Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Thr Glu  Pro Ser Glu
    1640                1645                1650

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1655                1660                1665

Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Ser
    1670                1675                1680

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Ser  Pro Ala Gly
    1685                1690                1695

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1700                1705                1710

Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    1715                1720                1725

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    1730                1735                1740

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Thr Glu
    1745                1750                1755

Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Thr Glu  Pro Ser Glu
    1760                1765                1770

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1775                1780                1785

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    1790                1795                1800

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    1805                1810                1815
```

```
Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Thr Glu  Pro Ser Glu
    1820                 1825                 1830

Gly Ser  Ala Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    1835                 1840                 1845

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1850                 1855                 1860

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Pro Ala Gly
    1865                 1870                 1875

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    1880                 1885                 1890

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1895                 1900                 1905

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    1910                 1915                 1920

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Thr  Ser Thr Glu
    1925                 1930                 1935

Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Thr Glu  Pro Ser Glu
    1940                 1945                 1950

Gly Ser  Ala Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    1955                 1960                 1965

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    1970                 1975                 1980

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    1985                 1990                 1995

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    2000                 2005                 2010

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2015                 2020                 2025

Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr
    2030                 2035                 2040

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    2045                 2050                 2055

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2060                 2065                 2070
```

```
Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    2075                 2080                 2085

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2090                 2095                 2100

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2105                 2110                 2115

Ser Pro  Thr Ser Thr Glu Glu  Gly Ser Pro Ala Gly  Ser Pro Thr
    2120                 2125                 2130

Ser Thr  Glu Glu Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2135                 2140                 2145

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    2150                 2155                 2160

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    2165                 2170                 2175

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2180                 2185                 2190

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    2195                 2200                 2205

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    2210                 2215                 2220

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly
    2225                 2230                 2235

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro
    2240                 2245                 2250

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    2255                 2260                 2265

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    2270                 2275                 2280

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly
    2285                 2290                 2295

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    2300                 2305                 2310

Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    2315                 2320                 2325
```

```
Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr
    2330                 2335                 2340

Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala
    2345                 2350                 2355

Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    2360                 2365                 2370

Ser Glu  Thr Pro Gly Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu
    2375                 2380                 2385

Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr
    2390                 2395                 2400

Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro Gly Ser  Glu Pro Ala
    2405                 2410                 2415

Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    2420                 2425                 2430

Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala
    2435                 2440                 2445

Pro
```

<210> 1540
<211> 2450
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1540

```
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1               5                  10                  15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
```

85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe

355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu

625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
740 745 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
755 760 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
770 775 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785 790 795 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
805 810 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
820 825 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
835 840 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
850 855 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865 870 875 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
885 890 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe

900 905 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
915 920 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
930 935 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945 950 955 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
965 970 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
980 985 990

Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
995 1000 1005

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1010 1015 1020

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1025 1030 1035

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1040 1045 1050

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1055 1060 1065

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1070 1075 1080

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1085 1090 1095

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1100 1105 1110

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1115 1120 1125

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1130 1135 1140

Thr Ala Ser Gly Gln Tyr Gly Pro Gly Ser Ser Pro Ser Ala Ser
1145 1150 1155

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr

```
    1160                1165                1170

Gly Pro  Gly Thr Pro Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly
    1175                1180                1185

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Pro
    1190                1195                1200

Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ala Ser Pro  Gly Thr Ser
    1205                1210                1215

Ser Thr  Gly Ser Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
    1220                1225                1230

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly
    1235                1240                1245

Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser Pro Gly  Ala Ser Pro
    1250                1255                1260

Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1265                1270                1275

Ser Thr  Gly Ser Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
    1280                1285                1290

Ser Gly  Gly Gln Trp Ala Pro  Lys Leu Ala Arg Leu  His Tyr Ser
    1295                1300                1305

Gly Ser  Ile Asn Ala Trp Ser  Thr Lys Glu Pro Phe  Ser Trp Ile
    1310                1315                1320

Lys Val  Asp Leu Leu Ala Pro  Met Ile Ile His Gly  Ile Lys Thr
    1325                1330                1335

Gln Gly  Ala Arg Gln Lys Phe  Ser Ser Leu Tyr Ile  Ser Gln Phe
    1340                1345                1350

Ile Ile  Met Tyr Ser Leu Asp  Gly Lys Lys Trp Gln  Thr Tyr Arg
    1355                1360                1365

Gly Asn  Ser Thr Gly Thr Leu  Met Val Phe Phe Gly  Asn Val Asp
    1370                1375                1380

Ser Ser  Gly Ile Lys His Asn  Ile Phe Asn Pro Pro  Ile Ile Ala
    1385                1390                1395

Arg Tyr  Ile Arg Leu His Pro  Thr His Tyr Ser Ile  Arg Ser Thr
    1400                1405                1410

Leu Arg  Met Glu Leu Met Gly  Cys Asp Leu Asn Ser  Cys Ser Met
```

1415 1420 1425

Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp Ala Gln Ile Thr
1430 1435 1440

Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp Ser Pro Ser
1445 1450 1455

Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg Pro
1460 1465 1470

Gln Gly Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
1475 1480 1485

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
1490 1495 1500

Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr
1505 1510 1515

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
1520 1525 1530

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
1535 1540 1545

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
1550 1555 1560

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
1565 1570 1575

Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
1580 1585 1590

Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
1595 1600 1605

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
1610 1615 1620

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
1625 1630 1635

Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
1640 1645 1650

Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
1655 1660 1665

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro

1670 1675 1680

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
1685 1690 1695

Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
1700 1705 1710

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
1715 1720 1725

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
1730 1735 1740

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
1745 1750 1755

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
1760 1765 1770

Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
1775 1780 1785

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
1790 1795 1800

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
1805 1810 1815

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
1820 1825 1830

Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
1835 1840 1845

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
1850 1855 1860

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly
1865 1870 1875

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
1880 1885 1890

Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
1895 1900 1905

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
1910 1915 1920

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly

1925 1930 1935

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
1940 1945 1950

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
1955 1960 1965

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
1970 1975 1980

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
1985 1990 1995

Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
2000 2005 2010

Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
2015 2020 2025

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
2030 2035 2040

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
2045 2050 2055

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
2060 2065 2070

Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
2075 2080 2085

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
2090 2095 2100

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
2105 2110 2115

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
2120 2125 2130

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
2135 2140 2145

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
2150 2155 2160

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
2165 2170 2175

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly

2180 2185 2190

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
2195 2200 2205

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
2210 2215 2220

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
2225 2230 2235

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
2240 2245 2250

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
2255 2260 2265

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
2270 2275 2280

Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr
2285 2290 2295

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
2300 2305 2310

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
2315 2320 2325

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Val Asn Asn
2330 2335 2340

Pro Lys Glu Trp Leu Gln Val Asp Phe Gln Lys Thr Met Lys Val
2345 2350 2355

Thr Gly Val Thr Thr Gln Gly Val Lys Ser Leu Leu Thr Ser Met
2360 2365 2370

Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly His Gln
2375 2380 2385

Trp Thr Leu Phe Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly
2390 2395 2400

Asn Gln Asp Ser Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro
2405 2410 2415

Leu Leu Thr Arg Tyr Leu Arg Ile His Pro Gln Ser Trp Val His
2420 2425 2430

Gln Ile Ala Leu Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp

2435 2440 2445

Leu Tyr
2450

<210> 1541
<211> 1715
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1541
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

```
His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480
```

```
Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750
```

```
Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Gly Gly Ser Glu Pro Ala Thr
            900                 905                 910

Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
        915                 920                 925

Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser
    930                 935                 940

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro
945                 950                 955                 960

Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu
                965                 970                 975

Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser
            980                 985                 990

Glu Pro Ala Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Thr Glu Pro
        995                 1000                1005

Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1010                1015                1020
```

```
Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro
    1025                1030                1035

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Ala Glu
    1040                1045                1050

Pro Arg  Lys Asn Phe Val Lys  Pro Asn Glu Thr Lys  Thr Tyr Phe
    1055                1060                1065

Trp Lys  Val Gln His His Met  Ala Pro Thr Lys Asp  Glu Phe Asp
    1070                1075                1080

Cys Lys  Ala Trp Ala Tyr Phe  Ser Asp Val Asp Leu  Glu Lys Asp
    1085                1090                1095

Val His  Ser Gly Leu Ile Gly  Pro Leu Leu Val Cys  His Thr Asn
    1100                1105                1110

Thr Leu  Asn Pro Ala His Gly  Arg Gln Val Thr Val  Gln Glu Phe
    1115                1120                1125

Ala Leu  Phe Phe Thr Ile Phe  Asp Glu Thr Lys Ser  Trp Tyr Phe
    1130                1135                1140

Thr Glu  Asn Met Glu Arg Asn  Cys Arg Ala Pro Cys  Asn Ile Gln
    1145                1150                1155

Met Glu  Asp Pro Thr Phe Lys  Glu Asn Tyr Arg Phe  His Ala Ile
    1160                1165                1170

Asn Gly  Tyr Ile Met Asp Thr  Leu Pro Gly Leu Val  Met Ala Gln
    1175                1180                1185

Asp Gln  Arg Ile Arg Trp Tyr  Leu Leu Ser Met Gly  Ser Asn Glu
    1190                1195                1200

Asn Ile  His Ser Ile His Phe  Ser Gly His Val Phe  Thr Val Arg
    1205                1210                1215

Lys Lys  Glu Glu Tyr Lys Met  Ala Leu Tyr Asn Leu  Tyr Pro Gly
    1220                1225                1230

Val Phe  Glu Thr Val Glu Met  Leu Pro Ser Lys Ala  Gly Ile Trp
    1235                1240                1245

Arg Val  Glu Cys Leu Ile Gly  Glu His Leu His Ala  Gly Met Ser
    1250                1255                1260

Thr Leu  Phe Leu Val Tyr Ser  Asn Lys Cys Gln Thr  Pro Leu Gly
    1265                1270                1275
```

```
Met Ala  Ser Gly His Ile Arg  Asp Phe Gln Ile Thr  Ala Ser Gly
    1280                1285                1290

Gln Tyr  Gly Gln Trp Ala Pro  Lys Leu Ala Arg Leu  His Tyr Ser
    1295                1300                1305

Gly Ser  Ile Asn Ala Trp Ser  Thr Lys Glu Pro Phe  Ser Trp Ile
    1310                1315                1320

Lys Val  Asp Leu Leu Ala Pro  Met Ile Ile His Gly  Ile Lys Thr
    1325                1330                1335

Gln Gly  Ala Arg Gln Lys Phe  Gly Ser Glu Pro Ala  Thr Ser Gly
    1340                1345                1350

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1355                1360                1365

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    1370                1375                1380

Pro Ala  Gly Ser Pro Gly Ser  Ser Leu Tyr Ile Ser  Gln Phe Ile
    1385                1390                1395

Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp Gln Thr  Tyr Arg Gly
    1400                1405                1410

Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe Gly Asn  Val Asp Ser
    1415                1420                1425

Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro Pro Ile  Ile Ala Arg
    1430                1435                1440

Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser Ile Arg  Ser Thr Leu
    1445                1450                1455

Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn Ser Cys  Ser Met Pro
    1460                1465                1470

Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp Ala Gln  Ile Thr Ala
    1475                1480                1485

Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr Trp Ser  Pro Ser Lys
    1490                1495                1500

Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn Ala Trp  Arg Pro Gln
    1505                1510                1515

Val Gly  Pro Ala Gly Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser
    1520                1525                1530
```

```
Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala
    1535                1540                1545

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Gly
    1550                1555                1560

Asn Asn  Pro Lys Glu Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met
    1565                1570                1575

Lys Val  Thr Gly Val Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr
    1580                1585                1590

Ser Met  Tyr Val Lys Glu Phe  Leu Ile Ser Ser Ser  Gln Asp Gly
    1595                1600                1605

His Gln  Trp Thr Leu Phe Phe  Gln Asn Gly Gly Thr  Glu Pro Ser
    1610                1615                1620

Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr
    1625                1630                1635

Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    1640                1645                1650

Ser Glu  Pro Ala Thr Ser Gly  Ser Lys Val Lys Val  Phe Gln Gly
    1655                1660                1665

Asn Gln  Asp Ser Phe Thr Pro  Val Val Asn Ser Leu  Asp Pro Pro
    1670                1675                1680

Leu Leu  Thr Arg Tyr Leu Arg  Ile His Pro Gln Ser  Trp Val His
    1685                1690                1695

Gln Ile  Ala Leu Arg Met Glu  Val Leu Gly Cys Glu  Ala Gln Asp
    1700                1705                1710

Leu Tyr
    1715
```

<210> 1542
<211> 3317
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1542

```
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1               5                  10                  15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Gly Pro Gly
```

```
            20                  25                  30

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser
        35                  40                  45

Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
    50                  55                  60

Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
65                  70                  75                  80

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly
                85                  90                  95

Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
            100                 105                 110

Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
        115                 120                 125

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly
    130                 135                 140

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
145                 150                 155                 160

Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Gly Phe
                165                 170                 175

Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr
            180                 185                 190

Lys Lys Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala
        195                 200                 205

Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala
    210                 215                 220

Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His
225                 230                 235                 240

Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu
                245                 250                 255

Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp
            260                 265                 270

Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys
        275                 280                 285

Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr
```

290 295 300

Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly
305 310 315 320

Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln
325 330 335

Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys
340 345 350

Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala
355 360 365

Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val
370 375 380

Asn Arg Ser Leu Pro Gly Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
385 390 395 400

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
405 410 415

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
420 425 430

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
435 440 445

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
450 455 460

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
465 470 475 480

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
485 490 495

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
500 505 510

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
515 520 525

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
530 535 540

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
545 550 555 560

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro

565 570 575

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
580 585 590

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
595 600 605

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
610 615 620

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
625 630 635 640

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
645 650 655

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
660 665 670

Pro Ser Gly Ala Thr Gly Ser Gly Leu Ile Gly Cys His Arg Lys Ser
675 680 685

Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser
690 695 700

Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala
705 710 715 720

Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu
725 730 735

Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln
740 745 750

His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu
755 760 765

Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp
770 775 780

Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn
785 790 795 800

Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys
805 810 815

Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala
820 825 830

Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu

835 840 845

Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe
850 855 860

Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His
865 870 875 880

Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr
885 890 895

Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr
900 905 910

Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro
915 920 925

Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile
930 935 940

Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser
945 950 955 960

Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu
965 970 975

Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys
980 985 990

Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn
995 1000 1005

Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu
1010 1015 1020

Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln
1025 1030 1035

Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile
1040 1045 1050

Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His
1055 1060 1065

Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp
1070 1075 1080

Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met
1085 1090 1095

Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr

1100 1105 1110

Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys
1115 1120 1125

His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys
1130 1135 1140

Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser
1145 1150 1155

Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile
1160 1165 1170

Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
1175 1180 1185

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu
1190 1195 1200

Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp
1205 1210 1215

Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe
1220 1225 1230

Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu
1235 1240 1245

Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg
1250 1255 1260

Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln
1265 1270 1275

Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu
1280 1285 1290

Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu
1295 1300 1305

Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg
1310 1315 1320

Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln
1325 1330 1335

Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu
1340 1345 1350

Thr Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr

1355 1360 1365

Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val
1370 1375 1380

Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu
1385 1390 1395

Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val
1400 1405 1410

Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
1415 1420 1425

Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala
1430 1435 1440

Pro Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr
1445 1450 1455

Arg Phe His Ala Gly Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro
1460 1465 1470

Gly Leu Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu
1475 1480 1485

Ser Met Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly
1490 1495 1500

His Val Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu
1505 1510 1515

Tyr Asn Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro
1520 1525 1530

Ser Lys Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His
1535 1540 1545

Leu His Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys
1550 1555 1560

Cys Gln Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe
1565 1570 1575

Gln Ile Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu
1580 1585 1590

Ala Arg Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys
1595 1600 1605

Glu Pro Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile

1610 1615 1620

Ile His Gly Ile Lys Thr Gln Gly Ala Arg Gly Ser Pro Ala Gly
1625 1630 1635

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
1640 1645 1650

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
1655 1660 1665

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr
1670 1675 1680

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
1685 1690 1695

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
1700 1705 1710

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
1715 1720 1725

Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser
1730 1735 1740

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
1745 1750 1755

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
1760 1765 1770

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
1775 1780 1785

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr
1790 1795 1800

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
1805 1810 1815

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
1820 1825 1830

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
1835 1840 1845

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
1850 1855 1860

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu

1865 1870 1875

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
1880 1885 1890

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1895 1900 1905

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
1910 1915 1920

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
1925 1930 1935

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
1940 1945 1950

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1955 1960 1965

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
1970 1975 1980

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
1985 1990 1995

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
2000 2005 2010

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2015 2020 2025

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
2030 2035 2040

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu
2045 2050 2055

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
2060 2065 2070

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr
2075 2080 2085

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
2090 2095 2100

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
2105 2110 2115

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu

2120 2125 2130

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
2135 2140 2145

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser
2150 2155 2160

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
2165 2170 2175

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
2180 2185 2190

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
2195 2200 2205

Pro Gly Gln Lys Phe Ser Ser Leu Tyr Ile Ser Gln Phe Ile Ile
2210 2215 2220

Met Tyr Ser Leu Asp Gly Lys Lys Trp Gln Thr Tyr Arg Gly Asn
2225 2230 2235

Ser Thr Gly Thr Leu Met Val Phe Phe Gly Asn Val Asp Ser Ser
2240 2245 2250

Gly Ile Lys His Asn Ile Phe Asn Pro Pro Ile Ile Ala Arg Tyr
2255 2260 2265

Ile Arg Leu His Pro Thr His Tyr Ser Ile Arg Ser Thr Leu Arg
2270 2275 2280

Met Glu Leu Met Gly Cys Asp Leu Asn Ser Cys Ser Met Pro Leu
2285 2290 2295

Gly Met Glu Ser Lys Ala Ile Ser Asp Ala Gln Ile Thr Ala Ser
2300 2305 2310

Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp Ser Pro Ser Lys Ala
2315 2320 2325

Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg Pro Gln Val
2330 2335 2340

Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln Lys Thr Met
2345 2350 2355

Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser Leu Leu Thr
2360 2365 2370

Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly

2375 2380 2385

His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val Lys Val Phe
2390 2395 2400

Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn Ser Leu Asp
2405 2410 2415

Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro Gln Ser Trp
2420 2425 2430

Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly Cys Glu Ala
2435 2440 2445

Gln Asp Leu Tyr Gly Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr
2450 2455 2460

Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
2465 2470 2475

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
2480 2485 2490

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
2495 2500 2505

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
2510 2515 2520

Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
2525 2530 2535

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
2540 2545 2550

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro
2555 2560 2565

Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
2570 2575 2580

Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
2585 2590 2595

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
2600 2605 2610

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
2615 2620 2625

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser

2630 2635 2640

Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
2645 2650 2655

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
2660 2665 2670

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
2675 2680 2685

Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
2690 2695 2700

Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
2705 2710 2715

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu
2720 2725 2730

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro
2735 2740 2745

Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
2750 2755 2760

Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
2765 2770 2775

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
2780 2785 2790

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
2795 2800 2805

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
2810 2815 2820

Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
2825 2830 2835

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
2840 2845 2850

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro
2855 2860 2865

Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
2870 2875 2880

Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly

2885 2890 2895

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
2900 2905 2910

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
2915 2920 2925

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
2930 2935 2940

Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
2945 2950 2955

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
2960 2965 2970

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro
2975 2980 2985

Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr
2990 2995 3000

Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
3005 3010 3015

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
3020 3025 3030

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
3035 3040 3045

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
3050 3055 3060

Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
3065 3070 3075

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
3080 3085 3090

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro
3095 3100 3105

Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
3110 3115 3120

Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
3125 3130 3135

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala

3140 3145 3150

Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro
3155 3160 3165

Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
3170 3175 3180

Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
3185 3190 3195

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu
3200 3205 3210

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
3215 3220 3225

Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu
3230 3235 3240

Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
3245 3250 3255

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
3260 3265 3270

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser
3275 3280 3285

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
3290 3295 3300

Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
3305 3310 3315

<210> 1543
<211> 3347
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1543
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

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Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320
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Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590
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Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Ser Arg His Pro Ser Thr Arg
            740                 745                 750

Gln Lys Gln Phe Asn Ala Thr Thr Ile Pro Glu Asn Asp Ile Glu Lys
        755                 760                 765

Thr Asp Pro Trp Phe Ala His Arg Thr Pro Met Pro Lys Ile Gln Asn
    770                 775                 780

Val Ser Ser Ser Asp Leu Leu Met Leu Leu Arg Gln Ser Pro Thr Pro
785                 790                 795                 800

His Gly Leu Ser Leu Ser Asp Leu Gln Glu Ala Lys Tyr Glu Thr Phe
                805                 810                 815

Ser Asp Asp Pro Ser Pro Gly Ala Ile Asp Ser Asn Asn Ser Leu Ser
            820                 825                 830

Glu Met Thr His Phe Arg Pro Gln Leu His His Ser Gly Asp Met Val
        835                 840                 845

Phe Thr Pro Glu Ser Gly Leu Gln Leu Arg Leu Asn Glu Lys Leu Gly
    850                 855                 860
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```
Thr Thr Ala Ala Thr Glu Leu Lys Lys Leu Asp Phe Lys Val Ser Ser
865                 870                 875                 880

Thr Ser Asn Asn Leu Ile Ser Thr Ile Pro Ser Asp Asn Leu Ala Ala
                885                 890                 895

Gly Thr Asp Asn Thr Ser Ser Leu Gly Pro Pro Ser Met Pro Val His
            900                 905                 910

Tyr Asp Ser Gln Leu Asp Thr Thr Leu Phe Gly Lys Lys Ser Ser Pro
        915                 920                 925

Leu Thr Glu Ser Gly Gly Pro Leu Ser Leu Ser Glu Glu Asn Asn Asp
    930                 935                 940

Ser Lys Leu Leu Glu Ser Gly Leu Met Asn Ser Gln Glu Ser Ser Trp
945                 950                 955                 960

Gly Lys Asn Val Ser Ser Thr Glu Ser Gly Arg Leu Phe Lys Gly Lys
                965                 970                 975

Arg Ala His Gly Pro Ala Leu Leu Thr Lys Asp Asn Ala Leu Phe Lys
            980                 985                 990

Val Ser Ile Ser Leu Leu Lys Thr  Asn Lys Thr Ser Asn  Asn Ser Ala
        995                 1000                1005

Thr Asn  Arg Lys Thr His Ile  Asp Gly Pro Ser Leu  Leu Ile Glu
    1010                1015                1020

Asn Ser  Pro Ser Val Trp Gln  Asn Ile Leu Glu Ser  Asp Thr Glu
    1025                1030                1035

Phe Lys  Lys Val Thr Pro Leu  Ile His Asp Arg Met  Leu Met Asp
    1040                1045                1050

Lys Asn  Ala Thr Ala Leu Arg  Leu Asn His Met Ser  Asn Lys Thr
    1055                1060                1065

Thr Ser  Ser Lys Asn Met Glu  Met Val Gln Gln Lys  Lys Glu Gly
    1070                1075                1080

Pro Ile  Pro Pro Asp Ala Gln  Asn Pro Asp Met Ser  Phe Phe Lys
    1085                1090                1095

Met Leu  Phe Leu Pro Glu Ser  Ala Arg Trp Ile Gln  Arg Thr His
    1100                1105                1110

Gly Lys  Asn Ser Leu Asn Ser  Gly Gln Gly Pro Ser  Pro Lys Gln
    1115                1120                1125
```

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Leu Val  Ser Leu Gly Pro Glu  Lys Ser Val Glu Gly  Gln Asn Phe
    1130                1135                1140

Leu Ser  Glu Lys Asn Lys Val  Val Val Gly Lys Gly  Glu Phe Thr
    1145                1150                1155

Lys Asp  Val Gly Leu Lys Glu  Met Val Phe Pro Ser  Ser Arg Asn
    1160                1165                1170

Leu Phe  Leu Thr Asn Leu Asp  Asn Leu His Glu Asn  Asn Thr His
    1175                1180                1185

Asn Gln  Glu Lys Lys Ile Gln  Glu Glu Ile Glu Lys  Lys Glu Thr
    1190                1195                1200

Leu Ile  Gln Glu Asn Val Val  Leu Pro Gln Ile His  Thr Val Thr
    1205                1210                1215

Gly Thr  Lys Asn Phe Met Lys  Asn Leu Phe Leu Leu  Ser Thr Arg
    1220                1225                1230

Gln Asn  Val Glu Gly Ser Tyr  Asp Gly Ala Tyr Ala  Pro Val Leu
    1235                1240                1245

Gln Asp  Phe Arg Ser Leu Asn  Asp Ser Thr Asn Arg  Thr Lys Lys
    1250                1255                1260

His Thr  Ala His Phe Ser Lys  Lys Gly Glu Glu Glu  Asn Leu Glu
    1265                1270                1275

Gly Leu  Gly Asn Gln Thr Lys  Gln Ile Val Glu Lys  Tyr Ala Cys
    1280                1285                1290

Thr Thr  Arg Ile Ser Pro Asn  Thr Ser Gln Gln Asn  Phe Val Thr
    1295                1300                1305

Gln Arg  Ser Lys Arg Ala Leu  Lys Gln Phe Arg Leu  Pro Leu Glu
    1310                1315                1320

Glu Thr  Glu Leu Glu Lys Arg  Ile Ile Val Asp Asp  Thr Ser Thr
    1325                1330                1335

Gln Trp  Ser Lys Asn Met Lys  His Leu Thr Pro Ser  Thr Leu Thr
    1340                1345                1350

Gln Ile  Asp Tyr Asn Glu Lys  Glu Lys Gly Ala Ile  Thr Gln Ser
    1355                1360                1365

Pro Leu  Ser Asp Cys Leu Thr  Arg Ser His Ser Ile  Pro Gln Ala
    1370                1375                1380
```

```
Asn Arg  Ser Pro Leu Pro Ile  Ala Lys Val Ser Ser  Phe Pro Ser
    1385                1390                1395

Ile Arg  Pro Ile Tyr Leu Thr  Arg Val Leu Phe Gln  Asp Asn Ser
    1400                1405                1410

Ser His  Leu Pro Ala Ala Ser  Tyr Arg Lys Lys Asp  Ser Gly Val
    1415                1420                1425

Gln Glu  Ser Ser His Phe Leu  Gln Gly Ala Lys Lys  Asn Asn Leu
    1430                1435                1440

Ser Leu  Ala Ile Leu Thr Leu  Glu Met Thr Gly Asp  Gln Arg Glu
    1445                1450                1455

Val Gly  Ser Leu Gly Thr Ser  Ala Thr Asn Ser Val  Thr Tyr Lys
    1460                1465                1470

Lys Val  Glu Asn Thr Val Leu  Pro Lys Pro Asp Leu  Pro Lys Thr
    1475                1480                1485

Ser Gly  Lys Val Glu Leu Leu  Pro Lys Val His Ile  Tyr Gln Lys
    1490                1495                1500

Asp Leu  Phe Pro Thr Glu Thr  Ser Asn Gly Ser Pro  Gly His Leu
    1505                1510                1515

Asp Leu  Val Glu Gly Ser Leu  Leu Gln Gly Thr Glu  Gly Ala Ile
    1520                1525                1530

Lys Trp  Asn Glu Ala Asn Arg  Pro Gly Lys Val Pro  Phe Leu Arg
    1535                1540                1545

Val Ala  Thr Glu Ser Ser Ala  Lys Thr Pro Ser Lys  Leu Leu Asp
    1550                1555                1560

Pro Leu  Ala Trp Asp Asn His  Tyr Gly Thr Gln Ile  Pro Lys Glu
    1565                1570                1575

Glu Trp  Lys Ser Gln Glu Lys  Ser Pro Glu Lys Thr  Ala Phe Lys
    1580                1585                1590

Lys Lys  Asp Thr Ile Leu Ser  Leu Asn Ala Cys Glu  Ser Asn His
    1595                1600                1605

Ala Ile  Ala Ala Ile Asn Glu  Gly Gln Asn Lys Pro  Glu Ile Glu
    1610                1615                1620

Val Thr  Trp Ala Lys Gln Gly  Arg Thr Glu Arg Leu  Cys Ser Gln
    1625                1630                1635
```

```
Asn Pro  Pro Val Leu Lys Arg  His Gln Arg Glu Ile  Thr Gly Pro
    1640                1645                1650

Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser Ser Pro  Gly Ser Ser
    1655                1660                1665

Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala
    1670                1675                1680

Ser Thr  Gly Thr Gly Pro Gly  Ser Ser Pro Ser Ala  Ser Thr Gly
    1685                1690                1695

Thr Gly  Pro Gly Ser Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro
    1700                1705                1710

Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Gly Ala Ser
    1715                1720                1725

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    1730                1735                1740

Ser Ser  Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
    1745                1750                1755

Gly Ser  Pro Gly Thr Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro
    1760                1765                1770

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1775                1780                1785

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    1790                1795                1800

Ser Ser  Thr Gly Ser Pro Gly  Ser Ser Pro Ser Ala  Ser Thr Gly
    1805                1810                1815

Thr Gly  Pro Gly Thr Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro
    1820                1825                1830

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1835                1840                1845

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    1850                1855                1860

Ser Ser  Thr Gly Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    1865                1870                1875

Gly Ser  Pro Gly Ser Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro
    1880                1885                1890
```

```
Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Thr Pro
    1895                 1900                 1905

Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly Ser Ser  Thr Pro Ser
    1910                 1915                 1920

Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    1925                 1930                 1935

Gly Ser  Pro Gly Ser Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro
    1940                 1945                 1950

Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr Gly Pro  Gly Ala Ser
    1955                 1960                 1965

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    1970                 1975                 1980

Ser Ser  Thr Gly Ser Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    1985                 1990                 1995

Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    2000                 2005                 2010

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    2015                 2020                 2025

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    2030                 2035                 2040

Ser Ser  Thr Gly Ser Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    2045                 2050                 2055

Ser Ser  Pro Gly Ser Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro
    2060                 2065                 2070

Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser Ser Pro  Gly Ser Ser
    2075                 2080                 2085

Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Thr Pro  Gly Ser Gly
    2090                 2095                 2100

Thr Ala  Ser Ser Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    2105                 2110                 2115

Gly Ser  Pro Gly Ser Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro
    2120                 2125                 2130

Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr Gly Pro  Gly Ser Ser
    2135                 2140                 2145
```

```
Pro Ser  Ala Ser Thr Gly Thr  Gly Pro Gly Ala Ser  Pro Gly Thr
    2150                2155                2160

Ser Ser  Thr Gly Ser Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    2165                2170                2175

Ser Ser  Pro Gly Ser Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro
    2180                2185                2190

Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr Gly Pro  Gly Ser Ser
    2195                2200                2205

Pro Ser  Ala Ser Thr Gly Thr  Gly Pro Gly Ala Ser  Pro Gly Thr
    2210                2215                2220

Ser Ser  Thr Gly Ser Ser Gly  Arg Thr Thr Leu Gln  Ser Asp Gln
    2225                2230                2235

Glu Glu  Ile Asp Tyr Asp Asp  Thr Ile Ser Val Glu  Met Lys Lys
    2240                2245                2250

Glu Asp  Phe Asp Ile Tyr Asp  Glu Asp Glu Asn Gln  Ser Pro Arg
    2255                2260                2265

Ser Phe  Gln Lys Lys Thr Arg  His Tyr Phe Ile Ala  Ala Val Glu
    2270                2275                2280

Arg Leu  Trp Asp Tyr Gly Met  Ser Ser Ser Pro His  Val Leu Arg
    2285                2290                2295

Asn Arg  Ala Gln Ser Gly Ser  Val Pro Gln Phe Lys  Lys Val Val
    2300                2305                2310

Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
    2315                2320                2325

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    2330                2335                2340

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    2345                2350                2355

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    2360                2365                2370

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Gly
    2375                2380                2385

Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser Ser Thr  Pro Ser Gly
    2390                2395                2400
```

```
Ala Thr  Gly Ser Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
    2405                 2410                 2415

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly
    2420                 2425                 2430

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Pro
    2435                 2440                 2445

Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ser Ser Pro  Ser Ala Ser
    2450                 2455                 2460

Thr Gly  Thr Gly Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    2465                 2470                 2475

Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly
    2480                 2485                 2490

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Pro
    2495                 2500                 2505

Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ser Ser Pro  Ser Ala Ser
    2510                 2515                 2520

Thr Gly  Thr Gly Pro Gly Ala  Ser Pro Gly Pro Asn  Glu Thr Lys
    2525                 2530                 2535

Thr Tyr  Phe Trp Lys Val Gln  His His Met Ala Pro  Thr Lys Asp
    2540                 2545                 2550

Glu Phe  Asp Cys Lys Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu
    2555                 2560                 2565

Glu Lys  Asp Val His Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys
    2570                 2575                 2580

His Thr  Asn Thr Leu Asn Pro  Ala His Gly Arg Gln  Val Thr Val
    2585                 2590                 2595

Gln Glu  Phe Ala Leu Phe Phe  Thr Ile Phe Asp Glu  Thr Lys Ser
    2600                 2605                 2610

Trp Tyr  Phe Thr Glu Asn Met  Glu Arg Asn Cys Arg  Ala Pro Cys
    2615                 2620                 2625

Asn Ile  Gln Met Glu Asp Pro  Thr Phe Lys Glu Asn  Tyr Arg Phe
    2630                 2635                 2640

His Ala  Ile Asn Gly Tyr Ile  Met Asp Thr Leu Pro  Gly Leu Val
    2645                 2650                 2655
```

```
Met Ala  Gln Asp Gln Arg Ile  Arg Trp Tyr Leu Leu  Ser Met Gly
    2660                 2665                 2670

Ser Asn  Glu Asn Ile His Ser  Ile His Phe Ser Gly  His Val Phe
    2675                 2680                 2685

Thr Val  Arg Lys Lys Glu Glu  Tyr Lys Met Ala Leu  Tyr Asn Leu
    2690                 2695                 2700

Tyr Pro  Gly Val Phe Glu Thr  Val Glu Met Leu Pro  Ser Lys Ala
    2705                 2710                 2715

Gly Ile  Trp Arg Val Glu Cys  Leu Ile Gly Glu His  Leu His Ala
    2720                 2725                 2730

Gly Met  Ser Thr Leu Phe Leu  Val Tyr Ser Asn Lys  Cys Gln Thr
    2735                 2740                 2745

Pro Leu  Gly Met Ala Ser Gly  His Ile Arg Asp Phe  Gln Ile Thr
    2750                 2755                 2760

Ala Ser  Gly Gln Tyr Gly Gln  Trp Ala Pro Lys Leu  Ala Arg Leu
    2765                 2770                 2775

His Tyr  Ser Gly Ser Ile Asn  Ala Trp Ser Thr Lys  Glu Pro Phe
    2780                 2785                 2790

Ser Trp  Ile Lys Val Asp Leu  Leu Ala Pro Met Ile  Ile His Gly
    2795                 2800                 2805

Ile Lys  Thr Gln Gly Ala Arg  Gln Lys Phe Gly Pro  Gly Ala Ser
    2810                 2815                 2820

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    2825                 2830                 2835

Ser Ser  Thr Gly Ser Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    2840                 2845                 2850

Ser Ser  Pro Gly Ser Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro
    2855                 2860                 2865

Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser Ser Pro  Gly Ser Ser
    2870                 2875                 2880

Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Thr Pro  Gly Ser Gly
    2885                 2890                 2895

Thr Ala  Ser Ser Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    2900                 2905                 2910
```

```
Gly Ser  Pro Gly Ser Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro
    2915                 2920                 2925

Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr Gly Pro  Gly Ser Ser
    2930                 2935                 2940

Pro Ser  Ala Ser Thr Gly Thr  Gly Pro Gly Ala Ser  Pro Gly Thr
    2945                 2950                 2955

Ser Ser  Thr Gly Ser Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    2960                 2965                 2970

Ser Ser  Pro Gly Ser Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro
    2975                 2980                 2985

Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr Gly Pro  Gly Ser Ser
    2990                 2995                 3000

Pro Ser  Ala Ser Thr Gly Thr  Gly Pro Gly Ala Ser  Pro Gly Thr
    3005                 3010                 3015

Ser Ser  Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
    3020                 3025                 3030

Gly Ser  Pro Gly Ser Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro
    3035                 3040                 3045

Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr Gly Pro  Gly Ala Ser
    3050                 3055                 3060

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala
    3065                 3070                 3075

Ser Thr  Gly Thr Gly Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    3080                 3085                 3090

Ser Ser  Pro Gly Ser Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Gly
    3095                 3100                 3105

Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile Met Tyr  Ser Leu Asp
    3110                 3115                 3120

Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn Ser Thr  Gly Thr Leu
    3125                 3130                 3135

Met Val  Phe Phe Gly Asn Val  Asp Ser Ser Gly Ile  Lys His Asn
    3140                 3145                 3150

Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr Ile Arg  Leu His Pro
    3155                 3160                 3165
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```
Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg Met Glu  Leu Met Gly
    3170                3175                3180

Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu Gly Met  Glu Ser Lys
    3185                3190                3195

Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser Ser Tyr  Phe Thr Asn
    3200                3205                3210

Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala Arg Leu  His Leu Gln
    3215                3220                3225

Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val Asn Asn  Pro Lys Glu
    3230                3235                3240

Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met Lys Val  Thr Gly Val
    3245                3250                3255

Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr Ser Met  Tyr Val Lys
    3260                3265                3270

Glu Phe  Leu Ile Ser Ser Ser  Gln Asp Gly His Gln  Trp Thr Leu
    3275                3280                3285

Phe Phe  Gln Asn Gly Lys Val  Lys Val Phe Gln Gly  Asn Gln Asp
    3290                3295                3300

Ser Phe  Thr Pro Val Val Asn  Ser Leu Asp Pro Pro  Leu Leu Thr
    3305                3310                3315

Arg Tyr  Leu Arg Ile His Pro  Gln Ser Trp Val His  Gln Ile Ala
    3320                3325                3330

Leu Arg  Met Glu Val Leu Gly  Cys Glu Ala Gln Asp  Leu Tyr
    3335                3340                3345


<210> 1544
<211> 1614
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
      polypeptide

<400> 1544
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1               5                  10                  15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Gly Gly Ala Pro
            20                  25                  30

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly
```

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        35                  40                  45

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
    50                  55                  60

Ser Pro Gly Pro Ser Gly Pro Gly Arg Phe Pro Pro Arg Val Pro Lys
65                  70                  75                  80

Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys Thr Leu Phe Val
                85                  90                  95

Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp
            100                 105                 110

Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val Tyr Asp Thr Val
        115                 120                 125

Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val Ser Leu His Ala
    130                 135                 140

Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp
145                 150                 155                 160

Gln Thr Ser Gln Arg Glu Lys Glu Gly Gly Pro Gly Thr Pro Gly Ser
                165                 170                 175

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
            180                 185                 190

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
        195                 200                 205

Ala Ser Pro Gly Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr
    210                 215                 220

Val Trp Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu
225                 230                 235                 240

Cys Leu Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu
                245                 250                 255

Asn Ser Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu
            260                 265                 270

Ala Lys Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala
        275                 280                 285

Val Phe Asp Glu Gly Gly Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
    290                 295                 300

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
```

305 310 315 320

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
325 330 335

Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg
340 345 350

Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly
355 360 365

Tyr Val Asn Ser Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser
370 375 380

Val Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser
385 390 395 400

Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala
405 410 415

Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu
420 425 430

Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln
435 440 445

His Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu
450 455 460

Pro Gly Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr
465 470 475 480

Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
485 490 495

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Gly Gln Leu Arg
500 505 510

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
515 520 525

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
530 535 540

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
545 550 555 560

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
565 570 575

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro

580 585 590

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
595 600 605

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
610 615 620

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
625 630 635 640

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
645 650 655

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
660 665 670

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
675 680 685

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
690 695 700

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
705 710 715 720

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
725 730 735

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
740 745 750

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
755 760 765

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
770 775 780

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
785 790 795 800

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
805 810 815

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
820 825 830

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
835 840 845

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp

```
    850                 855                 860

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
865                 870                 875                 880

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
                885                 890                 895

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
            900                 905                 910

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
        915                 920                 925

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
    930                 935                 940

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
945                 950                 955                 960

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
                965                 970                 975

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
            980                 985                 990

Met Ser Ser Ser Pro His Val Leu  Arg Asn Arg Ala Gln  Ser Gly Ser
        995                 1000                1005

Val Pro  Gln Phe Lys Lys Val  Val Phe Gln Glu Phe  Thr Asp Gly
    1010                1015                1020

Ser Phe  Thr Gln Pro Leu Tyr  Arg Gly Glu Leu Asn  Glu His Leu
    1025                1030                1035

Gly Leu  Leu Gly Pro Tyr Ile  Arg Ala Glu Val Glu  Asp Asn Ile
    1040                1045                1050

Met Val  Thr Phe Arg Asn Gln  Ala Ser Arg Pro Tyr  Ser Phe Tyr
    1055                1060                1065

Ser Ser  Leu Ile Ser Tyr Glu  Glu Asp Gln Arg Gln  Gly Ala Glu
    1070                1075                1080

Pro Arg  Lys Asn Phe Val Lys  Pro Asn Glu Thr Lys  Thr Tyr Phe
    1085                1090                1095

Trp Lys  Val Gln His His Met  Ala Pro Thr Lys Asp  Glu Phe Asp
    1100                1105                1110

Cys Lys  Ala Trp Ala Tyr Phe  Ser Asp Val Asp Leu  Glu Lys Asp
```

1115 1120 1125

Val His Ser Gly Leu Ile Gly Pro Leu Leu Val Cys His Thr Asn
1130 1135 1140

Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr Val Gln Glu Phe
1145 1150 1155

Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys Ser Trp Tyr Phe
1160 1165 1170

Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro Cys Asn Ile Gln
1175 1180 1185

Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg Phe His Ala Ile
1190 1195 1200

Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu Val Met Ala Gln
1205 1210 1215

Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met Gly Ser Asn Glu
1220 1225 1230

Asn Ile His Ser Ile His Phe Ser Gly His Val Phe Thr Val Arg
1235 1240 1245

Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn Leu Tyr Pro Gly
1250 1255 1260

Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys Ala Gly Ile Trp
1265 1270 1275

Arg Val Glu Cys Leu Ile Gly Glu His Leu His Ala Gly Met Ser
1280 1285 1290

Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln Thr Pro Leu Gly
1295 1300 1305

Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile Thr Ala Ser Gly
1310 1315 1320

Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg Leu His Tyr Ser
1325 1330 1335

Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro Phe Ser Trp Ile
1340 1345 1350

Lys Val Asp Leu Leu Ala Pro Met Ile Ile His Gly Ile Lys Thr
1355 1360 1365

Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr Ile Ser Gln Phe

1370 1375 1380

Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp Gln Thr Tyr Arg
1385 1390 1395

Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe Gly Asn Val Asp
1400 1405 1410

Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro Pro Ile Ile Ala
1415 1420 1425

Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser Ile Arg Ser Thr
1430 1435 1440

Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn Ser Cys Ser Met
1445 1450 1455

Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp Ala Gln Ile Thr
1460 1465 1470

Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp Thr Pro Ser
1475 1480 1485

Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg Pro
1490 1495 1500

Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln Lys
1505 1510 1515

Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser Leu
1520 1525 1530

Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser Gln
1535 1540 1545

Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val Lys
1550 1555 1560

Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn Ser
1565 1570 1575

Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro Gln
1580 1585 1590

Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly Cys
1595 1600 1605

Glu Ala Gln Asp Leu Tyr
1610

<210> 1545

<211> 2914
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1545

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Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1               5                  10                  15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240
```

```
Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Gly Gly Ser Glu Pro Ala Thr
            340                 345                 350

Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
        355                 360                 365

Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser
    370                 375                 380

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro
385                 390                 395                 400

Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu
                405                 410                 415

Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser
            420                 425                 430

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro
        435                 440                 445

Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
    450                 455                 460

Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr
465                 470                 475                 480

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Tyr Asp Asp Asp Leu
                485                 490                 495

Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro
            500                 505                 510
```

```
Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp
        515                 520                 525

Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu
    530                 535                 540

Val Leu Ala Pro Asp Asp Gly Gly Thr Ser Thr Pro Glu Ser Gly Ser
545                 550                 555                 560

Ala Ser Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly
                565                 570                 575

Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser Ser
            580                 585                 590

Thr Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly
        595                 600                 605

Thr Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly
    610                 615                 620

Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Thr Pro
625                 630                 635                 640

Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser
                645                 650                 655

Pro Gly Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly
            660                 665                 670

Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Pro Ser
        675                 680                 685

Gly Glu Ser Ser Thr Ala Pro Gly Arg Ser Tyr Lys Ser Gln Tyr Leu
    690                 695                 700

Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe
705                 710                 715                 720

Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His
                725                 730                 735

Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr
            740                 745                 750

Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr
        755                 760                 765

Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro
    770                 775                 780
```

```
Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile
785                 790                 795                 800

Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser
                805                 810                 815

Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu
            820                 825                 830

Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys
        835                 840                 845

Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn
    850                 855                 860

Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr
865                 870                 875                 880

Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu
                885                 890                 895

Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr
            900                 905                 910

Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr
        915                 920                 925

Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe
    930                 935                 940

Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu
945                 950                 955                 960

Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn
                965                 970                 975

Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg
            980                 985                 990

Gly Met Thr Ala Leu Leu Lys Val  Ser Ser Cys Asp Lys  Asn Thr Gly
        995                 1000                1005

Asp Tyr  Tyr Glu Asp Ser Tyr  Glu Asp Ile Ser Ala  Tyr Leu Leu
    1010                1015                1020

Ser Lys  Asn Asn Ala Ile Glu  Pro Arg Ser Phe Ser  Gln Asn Ser
    1025                1030                1035

Arg His  Pro Ser Thr Arg Gln  Lys Gln Phe Asn Ala  Thr Thr Ile
    1040                1045                1050
```

```
Pro Glu  Asn Asp Ile Glu Lys  Thr Asp Pro Trp Phe  Ala His Arg
    1055                 1060                 1065

Thr Pro  Met Pro Lys Ile Gln  Asn Val Ser Ser Ser  Asp Leu Leu
    1070                 1075                 1080

Met Leu  Leu Arg Gln Ser Pro  Thr Pro His Gly Leu  Ser Leu Ser
    1085                 1090                 1095

Asp Leu  Gln Glu Ala Lys Tyr  Glu Thr Phe Ser Asp  Asp Pro Ser
    1100                 1105                 1110

Pro Gly  Ala Ile Asp Ser Asn  Asn Ser Leu Ser Glu  Met Thr His
    1115                 1120                 1125

Phe Arg  Pro Gln Leu His His  Ser Gly Asp Met Val  Phe Thr Pro
    1130                 1135                 1140

Glu Ser  Gly Leu Gln Leu Arg  Leu Asn Glu Lys Leu  Gly Thr Thr
    1145                 1150                 1155

Ala Ala  Thr Glu Leu Lys Lys  Leu Asp Phe Lys Val  Ser Ser Thr
    1160                 1165                 1170

Ser Asn  Asn Leu Ile Ser Thr  Ile Pro Ser Asp Asn  Leu Ala Ala
    1175                 1180                 1185

Gly Thr  Asp Asn Thr Ser Ser  Leu Gly Pro Pro Ser  Met Pro Val
    1190                 1195                 1200

His Tyr  Asp Ser Gln Leu Asp  Thr Thr Leu Phe Gly  Lys Lys Ser
    1205                 1210                 1215

Ser Pro  Leu Thr Glu Ser Gly  Gly Pro Leu Ser Leu  Ser Glu Glu
    1220                 1225                 1230

Asn Asn  Asp Ser Lys Leu Leu  Glu Ser Gly Leu Met  Asn Ser Gln
    1235                 1240                 1245

Glu Ser  Ser Trp Gly Lys Asn  Val Ser Ser Thr Glu  Ser Gly Arg
    1250                 1255                 1260

Leu Phe  Lys Gly Lys Arg Ala  His Gly Pro Ala Leu  Leu Thr Lys
    1265                 1270                 1275

Asp Asn  Ala Leu Phe Lys Val  Ser Ile Ser Leu Leu  Lys Thr Asn
    1280                 1285                 1290

Lys Thr  Ser Asn Asn Ser Ala  Thr Asn Arg Lys Thr  His Ile Asp
    1295                 1300                 1305
```

```
Gly Pro  Ser Leu Leu Ile Glu  Asn Ser Pro Ser Val  Trp Gln Asn
    1310                 1315                 1320

Ile Leu  Glu Ser Asp Thr Glu  Phe Lys Lys Val Thr  Pro Leu Ile
    1325                 1330                 1335

His Asp  Arg Met Leu Met Asp  Lys Asn Ala Thr Ala  Leu Arg Leu
    1340                 1345                 1350

Asn His  Met Ser Asn Lys Thr  Thr Ser Ser Lys Asn  Met Glu Met
    1355                 1360                 1365

Val Gln  Gln Lys Lys Glu Gly  Pro Ile Pro Pro Asp  Ala Gln Asn
    1370                 1375                 1380

Pro Asp  Met Ser Phe Phe Lys  Met Leu Phe Leu Pro  Glu Ser Ala
    1385                 1390                 1395

Arg Trp  Ile Gln Arg Thr His  Gly Lys Asn Ser Leu  Asn Ser Gly
    1400                 1405                 1410

Gln Gly  Pro Ser Pro Lys Gln  Leu Val Ser Leu Gly  Pro Glu Lys
    1415                 1420                 1425

Ser Val  Glu Gly Gln Asn Phe  Leu Ser Glu Lys Asn  Lys Val Val
    1430                 1435                 1440

Val Gly  Lys Gly Glu Phe Thr  Lys Asp Val Gly Leu  Lys Glu Met
    1445                 1450                 1455

Val Phe  Pro Ser Ser Arg Asn  Leu Phe Leu Thr Asn  Leu Asp Asn
    1460                 1465                 1470

Leu His  Glu Asn Asn Thr His  Asn Gln Glu Lys Lys  Ile Gln Glu
    1475                 1480                 1485

Glu Ile  Glu Lys Lys Glu Thr  Leu Ile Gln Glu Asn  Val Val Leu
    1490                 1495                 1500

Pro Gln  Ile His Thr Val Thr  Gly Thr Lys Asn Phe  Met Lys Asn
    1505                 1510                 1515

Leu Phe  Leu Leu Ser Thr Arg  Gln Asn Val Glu Gly  Ser Tyr Asp
    1520                 1525                 1530

Gly Ala  Tyr Ala Pro Val Leu  Gln Asp Phe Arg Ser  Leu Asn Asp
    1535                 1540                 1545

Ser Thr  Asn Arg Thr Lys Lys  His Thr Ala His Phe  Ser Lys Lys
    1550                 1555                 1560
```

```
Gly Glu  Glu Glu Asn Leu Glu  Gly Leu Gly Asn Gln  Thr Lys Gln
    1565                1570                1575

Ile Val  Glu Lys Tyr Ala Cys  Thr Thr Arg Ile Ser  Pro Asn Thr
    1580                1585                1590

Ser Gln  Gln Asn Phe Val Thr  Gln Arg Ser Lys Arg  Ala Leu Lys
    1595                1600                1605

Gln Phe  Arg Leu Pro Leu Glu  Glu Thr Glu Leu Glu  Lys Arg Ile
    1610                1615                1620

Ile Val  Asp Asp Thr Ser Thr  Gln Trp Ser Lys Asn  Met Lys His
    1625                1630                1635

Leu Thr  Pro Ser Thr Leu Thr  Gln Ile Asp Tyr Asn  Glu Lys Glu
    1640                1645                1650

Lys Gly  Ala Ile Thr Gln Ser  Pro Leu Ser Asp Cys  Leu Thr Arg
    1655                1660                1665

Ser His  Ser Ile Pro Gln Ala  Asn Arg Ser Pro Leu  Pro Ile Ala
    1670                1675                1680

Lys Val  Ser Ser Phe Pro Ser  Ile Arg Pro Ile Tyr  Leu Thr Arg
    1685                1690                1695

Val Leu  Phe Gln Asp Asn Ser  Ser His Leu Pro Ala  Ala Ser Tyr
    1700                1705                1710

Arg Lys  Lys Asp Ser Gly Val  Gln Glu Ser Ser His  Phe Leu Gln
    1715                1720                1725

Gly Ala  Lys Lys Asn Asn Leu  Ser Leu Ala Ile Leu  Thr Leu Glu
    1730                1735                1740

Met Thr  Gly Asp Gln Arg Glu  Val Gly Ser Leu Gly  Thr Ser Ala
    1745                1750                1755

Thr Asn  Ser Val Thr Tyr Lys  Lys Val Glu Asn Thr  Val Leu Pro
    1760                1765                1770

Lys Pro  Asp Leu Pro Lys Thr  Ser Gly Lys Val Glu  Leu Leu Pro
    1775                1780                1785

Lys Val  His Ile Tyr Gln Lys  Asp Leu Phe Pro Thr  Glu Thr Ser
    1790                1795                1800

Asn Gly  Ser Pro Gly His Leu  Asp Leu Val Glu Gly  Ser Leu Leu
    1805                1810                1815
```

```
Gln Gly  Thr Glu Gly Ala Ile  Lys Trp Asn Glu Ala  Asn Arg Pro
    1820                1825                1830

Gly Lys  Val Pro Phe Leu Arg  Val Ala Thr Glu Ser  Ser Ala Lys
    1835                1840                1845

Thr Pro  Ser Lys Leu Leu Asp  Pro Leu Ala Trp Asp  Asn His Tyr
    1850                1855                1860

Gly Thr  Gln Ile Pro Lys Glu  Glu Trp Lys Ser Gln  Glu Lys Ser
    1865                1870                1875

Pro Glu  Lys Thr Ala Phe Lys  Lys Lys Asp Thr Ile  Leu Ser Leu
    1880                1885                1890

Asn Ala  Cys Glu Ser Asn His  Ala Ile Ala Ala Ile  Asn Glu Gly
    1895                1900                1905

Gln Asn  Lys Pro Glu Ile Glu  Val Thr Trp Ala Lys  Gln Gly Arg
    1910                1915                1920

Thr Glu  Arg Leu Cys Ser Gln  Asn Pro Pro Val Leu  Lys Arg His
    1925                1930                1935

Gln Arg  Glu Ile Thr Arg Thr  Thr Leu Gln Ser Asp  Gln Glu Glu
    1940                1945                1950

Ile Asp  Tyr Asp Asp Thr Ile  Ser Val Glu Met Lys  Lys Glu Asp
    1955                1960                1965

Phe Asp  Ile Tyr Asp Glu Asp  Glu Asn Gln Ser Pro  Arg Ser Phe
    1970                1975                1980

Gln Lys  Lys Thr Arg His Tyr  Phe Ile Ala Ala Val  Glu Arg Leu
    1985                1990                1995

Trp Asp  Tyr Gly Met Ser Ser  Ser Pro His Val Leu  Arg Asn Arg
    2000                2005                2010

Ala Gln  Ser Gly Ser Val Pro  Gln Phe Lys Lys Val  Val Phe Gln
    2015                2020                2025

Glu Phe  Thr Asp Gly Ser Phe  Thr Gln Pro Leu Tyr  Arg Gly Glu
    2030                2035                2040

Leu Asn  Glu His Leu Gly Leu  Leu Gly Pro Tyr Ile  Arg Ala Glu
    2045                2050                2055

Val Glu  Asp Asn Ile Met Val  Thr Phe Arg Asn Gln  Ala Ser Arg
    2060                2065                2070
```

```
Pro Tyr  Ser Phe Tyr Ser Ser  Leu Ile Ser Tyr Glu  Glu Asp Gln
    2075                2080                2085

Arg Gly  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    2090                2095                2100

Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr Pro Gly  Thr Ser Glu
    2105                2110                2115

Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser
    2120                2125                2130

Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    2135                2140                2145

Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    2150                2155                2160

Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Glu
    2165                2170                2175

Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser
    2180                2185                2190

Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    2195                2200                2205

Gly Pro  Gly Ser Pro Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly
    2210                2215                2220

Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr
    2225                2230                2235

Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr Ser Glu  Ser Ala Thr
    2240                2245                2250

Pro Glu  Ser Gly Pro Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    2255                2260                2265

Gly Pro  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    2270                2275                2280

Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr Pro Gly  Ser Glu Pro
    2285                2290                2295

Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Ser Pro Ala  Gly Ser Pro
    2300                2305                2310

Thr Ser  Thr Glu Glu Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    2315                2320                2325
```

```
Ala Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    2330                 2335                 2340

Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr Pro Gly  Thr Ser Glu
    2345                 2350                 2355

Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser
    2360                 2365                 2370

Glu Gly  Ser Ala Pro Gly Gln  Gly Ala Glu Pro Arg  Lys Asn Phe
    2375                 2380                 2385

Val Lys  Pro Asn Glu Thr Lys  Thr Tyr Phe Trp Lys  Val Gln His
    2390                 2395                 2400

His Met  Ala Pro Thr Lys Asp  Glu Phe Asp Cys Lys  Ala Trp Ala
    2405                 2410                 2415

Tyr Phe  Ser Asp Val Asp Leu  Glu Lys Asp Val His  Ser Gly Leu
    2420                 2425                 2430

Ile Gly  Pro Leu Leu Val Cys  His Thr Asn Thr Leu  Asn Pro Ala
    2435                 2440                 2445

His Gly  Arg Gln Val Thr Val  Gln Glu Phe Ala Leu  Phe Phe Thr
    2450                 2455                 2460

Ile Phe  Asp Glu Thr Lys Ser  Trp Tyr Phe Thr Glu  Asn Met Glu
    2465                 2470                 2475

Arg Asn  Cys Arg Ala Pro Cys  Asn Ile Gln Met Glu  Asp Pro Thr
    2480                 2485                 2490

Phe Lys  Glu Asn Tyr Arg Phe  His Ala Ile Asn Gly  Tyr Ile Met
    2495                 2500                 2505

Asp Thr  Leu Pro Gly Leu Val  Met Ala Gln Asp Gln  Arg Ile Arg
    2510                 2515                 2520

Trp Tyr  Leu Leu Ser Met Gly  Ser Asn Glu Asn Ile  His Ser Ile
    2525                 2530                 2535

His Phe  Ser Gly His Val Phe  Thr Val Arg Lys Lys  Glu Glu Tyr
    2540                 2545                 2550

Lys Met  Ala Leu Tyr Asn Leu  Tyr Pro Gly Val Phe  Glu Thr Val
    2555                 2560                 2565

Glu Met  Leu Pro Ser Lys Ala  Gly Ile Trp Arg Val  Glu Cys Leu
    2570                 2575                 2580
```

```
Ile Gly  Glu His Leu His Ala  Gly Met Ser Thr Leu  Phe Leu Val
    2585                 2590                 2595

Tyr Ser  Asn Lys Cys Gln Thr  Pro Leu Gly Met Ala  Ser Gly His
    2600                 2605                 2610

Ile Arg  Asp Phe Gln Ile Thr  Ala Ser Gly Gln Tyr  Gly Gln Trp
    2615                 2620                 2625

Ala Pro  Lys Leu Ala Arg Leu  His Tyr Ser Gly Ser  Ile Asn Ala
    2630                 2635                 2640

Trp Ser  Thr Lys Glu Pro Phe  Ser Trp Ile Lys Val  Asp Leu Leu
    2645                 2650                 2655

Ala Pro  Met Ile Ile His Gly  Ile Lys Thr Gln Gly  Ala Arg Gln
    2660                 2665                 2670

Lys Phe  Ser Ser Leu Tyr Ile  Ser Gln Phe Ile Ile  Met Tyr Ser
    2675                 2680                 2685

Leu Asp  Gly Lys Lys Trp Gln  Thr Tyr Arg Gly Asn  Ser Thr Gly
    2690                 2695                 2700

Thr Leu  Met Val Phe Phe Gly  Asn Val Asp Ser Ser  Gly Ile Lys
    2705                 2710                 2715

His Asn  Ile Phe Asn Pro Pro  Ile Ile Ala Arg Tyr  Ile Arg Leu
    2720                 2725                 2730

His Pro  Thr His Tyr Ser Ile  Arg Ser Thr Leu Arg  Met Glu Leu
    2735                 2740                 2745

Met Gly  Cys Asp Leu Asn Ser  Cys Ser Met Pro Leu  Gly Met Glu
    2750                 2755                 2760

Ser Lys  Ala Ile Ser Asp Ala  Gln Ile Thr Ala Ser  Ser Tyr Phe
    2765                 2770                 2775

Thr Asn  Met Phe Ala Thr Trp  Ser Pro Ser Lys Ala  Arg Leu His
    2780                 2785                 2790

Leu Gln  Gly Arg Ser Asn Ala  Trp Arg Pro Gln Val  Asn Asn Pro
    2795                 2800                 2805

Lys Glu  Trp Leu Gln Val Asp  Phe Gln Lys Thr Met  Lys Val Thr
    2810                 2815                 2820

Gly Val  Thr Thr Gln Gly Val  Lys Ser Leu Leu Thr  Ser Met Tyr
    2825                 2830                 2835
```

Val Lys Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly His Gln Trp
2840 2845 2850

Thr Leu Phe Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn
2855 2860 2865

Gln Asp Ser Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu
2870 2875 2880

Leu Thr Arg Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln
2885 2890 2895

Ile Ala Leu Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu
2900 2905 2910

Tyr

<210> 1546
<211> 2304
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1546
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn

130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro

405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp

675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Gly Ser Pro Ala Gly Ser
740 745 750

Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
755 760 765

Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
770 775 780

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro
785 790 795 800

Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
805 810 815

Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
820 825 830

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr
835 840 845

Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr
850 855 860

Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
865 870 875 880

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro
885 890 895

Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr
900 905 910

Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr
915 920 925

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala
930 935 940

Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser

945 950 955 960

Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
965 970 975

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro
980 985 990

Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
995 1000 1005

Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
1010 1015 1020

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala
1025 1030 1035

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr
1040 1045 1050

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu
1055 1060 1065

Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
1070 1075 1080

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
1085 1090 1095

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
1100 1105 1110

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
1115 1120 1125

Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
1130 1135 1140

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
1145 1150 1155

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
1160 1165 1170

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
1175 1180 1185

Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
1190 1195 1200

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro

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    1205                1210                1215

Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr
    1220                1225                1230

Pro Glu  Ser Gly Pro Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    1235                1240                1245

Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    1250                1255                1260

Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala
    1265                1270                1275

Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro
    1280                1285                1290

Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1295                1300                1305

Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    1310                1315                1320

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro
    1325                1330                1335

Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr
    1340                1345                1350

Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu
    1355                1360                1365

Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    1370                1375                1380

Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala Pro Gly  Ser Pro Ala
    1385                1390                1395

Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr Ser Glu  Ser Ala Thr
    1400                1405                1410

Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu
    1415                1420                1425

Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    1430                1435                1440

Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala
    1445                1450                1455

Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr Ser Thr  Glu Pro Ser
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1460 1465 1470

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
1475 1480 1485

Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
1490 1495 1500

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
1505 1510 1515

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser
1520 1525 1530

Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr
1535 1540 1545

Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
1550 1555 1560

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
1565 1570 1575

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
1580 1585 1590

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
1595 1600 1605

Ala Pro Gly Pro Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr
1610 1615 1620

Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp
1625 1630 1635

Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp
1640 1645 1650

Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg
1655 1660 1665

His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met
1670 1675 1680

Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
1685 1690 1695

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly
1700 1705 1710

Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu

1715 1720 1725

Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile
1730 1735 1740

Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr
1745 1750 1755

Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu
1760 1765 1770

Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe
1775 1780 1785

Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp Glu Phe Asp
1790 1795 1800

Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu Lys Asp
1805 1810 1815

Val His Ser Gly Leu Ile Gly Pro Leu Leu Val Cys His Thr Asn
1820 1825 1830

Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr Val Gln Glu Phe
1835 1840 1845

Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys Ser Trp Tyr Phe
1850 1855 1860

Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro Cys Asn Ile Gln
1865 1870 1875

Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg Phe His Ala Ile
1880 1885 1890

Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu Val Met Ala Gln
1895 1900 1905

Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met Gly Ser Asn Glu
1910 1915 1920

Asn Ile His Ser Ile His Phe Ser Gly His Val Phe Thr Val Arg
1925 1930 1935

Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn Leu Tyr Pro Gly
1940 1945 1950

Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys Ala Gly Ile Trp
1955 1960 1965

Arg Val Glu Cys Leu Ile Gly Glu His Leu His Ala Gly Met Ser

1970 1975 1980

Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln Thr Pro Leu Gly
1985 1990 1995

Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile Thr Ala Ser Gly
2000 2005 2010

Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg Leu His Tyr Ser
2015 2020 2025

Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro Phe Ser Trp Ile
2030 2035 2040

Lys Val Asp Leu Leu Ala Pro Met Ile Ile His Gly Ile Lys Thr
2045 2050 2055

Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr Ile Ser Gln Phe
2060 2065 2070

Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp Gln Thr Tyr Arg
2075 2080 2085

Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe Gly Asn Val Asp
2090 2095 2100

Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro Pro Ile Ile Ala
2105 2110 2115

Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser Ile Arg Ser Thr
2120 2125 2130

Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn Ser Cys Ser Met
2135 2140 2145

Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp Ala Gln Ile Thr
2150 2155 2160

Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp Ser Pro Ser
2165 2170 2175

Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg Pro
2180 2185 2190

Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln Lys
2195 2200 2205

Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser Leu
2210 2215 2220

Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser Gln

2225 2230 2235

Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val Lys
2240 2245 2250

Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn Ser
2255 2260 2265

Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro Gln
2270 2275 2280

Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly Cys
2285 2290 2295

Glu Ala Gln Asp Leu Tyr
2300

<210> 1547
<211> 1728
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1547
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

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Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415
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Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685
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Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Gly Thr Ser Glu Ser Ala
            740                 745                 750

Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu
        755                 760                 765

Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
    770                 775                 780

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala
785                 790                 795                 800

Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
                805                 810                 815

Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr
            820                 825                 830

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr
        835                 840                 845

Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
    850                 855                 860

Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser
865                 870                 875                 880

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro
                885                 890                 895

Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
            900                 905                 910

Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
        915                 920                 925

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr
    930                 935                 940

Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu
945                 950                 955                 960
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Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr
               965                 970                 975

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro
           980                 985                 990

Ser Glu Gly Ser Ala Pro Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu
       995                 1000                1005

Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
   1010                1015                1020

Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala Pro Gly  Pro Pro Val
   1025                1030                1035

Leu Lys  Arg His Gln Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Ser
   1040                1045                1050

Asp Gln  Glu Glu Ile Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met
   1055                1060                1065

Lys Lys  Glu Asp Phe Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser
   1070                1075                1080

Pro Arg  Ser Phe Gln Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala
   1085                1090                1095

Val Glu  Arg Leu Trp Asp Tyr  Gly Met Ser Ser Ser  Pro His Val
   1100                1105                1110

Leu Arg  Asn Arg Ala Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys
   1115                1120                1125

Val Val  Phe Gln Glu Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu
   1130                1135                1140

Tyr Arg  Gly Glu Leu Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr
   1145                1150                1155

Ile Arg  Ala Glu Val Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn
   1160                1165                1170

Gln Ala  Ser Arg Pro Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr
   1175                1180                1185

Glu Glu  Asp Gln Arg Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val
   1190                1195                1200

Lys Pro  Asn Glu Thr Lys Thr  Tyr Phe Trp Lys Val  Gln His His
   1205                1210                1215
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Met Ala  Pro Thr Lys Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr
    1220                1225                1230

Phe Ser  Asp Val Asp Leu Glu  Lys Asp Val His Ser  Gly Leu Ile
    1235                1240                1245

Gly Pro  Leu Leu Val Cys His  Thr Asn Thr Leu Asn  Pro Ala His
    1250                1255                1260

Gly Arg  Gln Val Thr Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile
    1265                1270                1275

Phe Asp  Glu Thr Lys Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg
    1280                1285                1290

Asn Cys  Arg Ala Pro Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe
    1295                1300                1305

Lys Glu  Asn Tyr Arg Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp
    1310                1315                1320

Thr Leu  Pro Gly Leu Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp
    1325                1330                1335

Tyr Leu  Leu Ser Met Gly Ser  Asn Glu Asn Ile His  Ser Ile His
    1340                1345                1350

Phe Ser  Gly His Val Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys
    1355                1360                1365

Met Ala  Leu Tyr Asn Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu
    1370                1375                1380

Met Leu  Pro Ser Lys Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile
    1385                1390                1395

Gly Glu  His Leu His Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr
    1400                1405                1410

Ser Asn  Lys Cys Gln Thr Pro  Leu Gly Met Ala Ser  Gly His Ile
    1415                1420                1425

Arg Asp  Phe Gln Ile Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala
    1430                1435                1440

Pro Lys  Leu Ala Arg Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp
    1445                1450                1455

Ser Thr  Lys Glu Pro Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala
    1460                1465                1470
```

```
Pro Met  Ile Ile His Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys
    1475                 1480                 1485

Phe Ser  Ser Leu Tyr Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu
    1490                 1495                 1500

Asp Gly  Lys Lys Trp Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr
    1505                 1510                 1515

Leu Met  Val Phe Phe Gly Asn  Val Asp Ser Ser Gly  Ile Lys His
    1520                 1525                 1530

Asn Ile  Phe Asn Pro Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His
    1535                 1540                 1545

Pro Thr  His Tyr Ser Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met
    1550                 1555                 1560

Gly Cys  Asp Leu Asn Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser
    1565                 1570                 1575

Lys Ala  Ile Ser Asp Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr
    1580                 1585                 1590

Asn Met  Phe Ala Thr Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu
    1595                 1600                 1605

Gln Gly  Arg Ser Asn Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys
    1610                 1615                 1620

Glu Trp  Leu Gln Val Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly
    1625                 1630                 1635

Val Thr  Thr Gln Gly Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val
    1640                 1645                 1650

Lys Glu  Phe Leu Ile Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr
    1655                 1660                 1665

Leu Phe  Phe Gln Asn Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln
    1670                 1675                 1680

Asp Ser  Phe Thr Pro Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu
    1685                 1690                 1695

Thr Arg  Tyr Leu Arg Ile His  Pro Gln Ser Trp Val  His Gln Ile
    1700                 1705                 1710

Ala Leu  Arg Met Glu Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr
    1715                 1720                 1725
```

<210> 1548
<211> 1728
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1548
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg

225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys

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            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gly Gly Thr Ser Glu Ser Ala Thr Pro
            740                 745                 750

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
        755                 760                 765

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
```

770 775 780

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
785 790 795 800

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
805 810 815

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
820 825 830

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
835 840 845

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
850 855 860

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala
865 870 875 880

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
885 890 895

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
900 905 910

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu
915 920 925

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
930 935 940

Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
945 950 955 960

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
965 970 975

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
980 985 990

Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
995 1000 1005

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
1010 1015 1020

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Gln Asn Pro Pro Val
1025 1030 1035

Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser

1040 1045 1050

Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
1055 1060 1065

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser
1070 1075 1080

Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala
1085 1090 1095

Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val
1100 1105 1110

Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys
1115 1120 1125

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu
1130 1135 1140

Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr
1145 1150 1155

Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn
1160 1165 1170

Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr
1175 1180 1185

Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val
1190 1195 1200

Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
1205 1210 1215

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr
1220 1225 1230

Phe Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile
1235 1240 1245

Gly Pro Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His
1250 1255 1260

Gly Arg Gln Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile
1265 1270 1275

Phe Asp Glu Thr Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg
1280 1285 1290

Asn Cys Arg Ala Pro Cys Asn Ile Gln Met Glu Asp Pro Thr Phe

1295 1300 1305

Lys Glu Asn Tyr Arg Phe His Ala Ile Asn Gly Tyr Ile Met Asp
1310 1315 1320

Thr Leu Pro Gly Leu Val Met Ala Gln Asp Gln Arg Ile Arg Trp
1325 1330 1335

Tyr Leu Leu Ser Met Gly Ser Asn Glu Asn Ile His Ser Ile His
1340 1345 1350

Phe Ser Gly His Val Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys
1355 1360 1365

Met Ala Leu Tyr Asn Leu Tyr Pro Gly Val Phe Glu Thr Val Glu
1370 1375 1380

Met Leu Pro Ser Lys Ala Gly Ile Trp Arg Val Glu Cys Leu Ile
1385 1390 1395

Gly Glu His Leu His Ala Gly Met Ser Thr Leu Phe Leu Val Tyr
1400 1405 1410

Ser Asn Lys Cys Gln Thr Pro Leu Gly Met Ala Ser Gly His Ile
1415 1420 1425

Arg Asp Phe Gln Ile Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala
1430 1435 1440

Pro Lys Leu Ala Arg Leu His Tyr Ser Gly Ser Ile Asn Ala Trp
1445 1450 1455

Ser Thr Lys Glu Pro Phe Ser Trp Ile Lys Val Asp Leu Leu Ala
1460 1465 1470

Pro Met Ile Ile His Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys
1475 1480 1485

Phe Ser Ser Leu Tyr Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu
1490 1495 1500

Asp Gly Lys Lys Trp Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr
1505 1510 1515

Leu Met Val Phe Phe Gly Asn Val Asp Ser Ser Gly Ile Lys His
1520 1525 1530

Asn Ile Phe Asn Pro Pro Ile Ile Ala Arg Tyr Ile Arg Leu His
1535 1540 1545

Pro Thr His Tyr Ser Ile Arg Ser Thr Leu Arg Met Glu Leu Met

1550 1555 1560

Gly Cys Asp Leu Asn Ser Cys Ser Met Pro Leu Gly Met Glu Ser
1565 1570 1575

Lys Ala Ile Ser Asp Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr
1580 1585 1590

Asn Met Phe Ala Thr Trp Ser Pro Ser Lys Ala Arg Leu His Leu
1595 1600 1605

Gln Gly Arg Ser Asn Ala Trp Arg Pro Gln Val Asn Asn Pro Lys
1610 1615 1620

Glu Trp Leu Gln Val Asp Phe Gln Lys Thr Met Lys Val Thr Gly
1625 1630 1635

Val Thr Thr Gln Gly Val Lys Ser Leu Leu Thr Ser Met Tyr Val
1640 1645 1650

Lys Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly His Gln Trp Thr
1655 1660 1665

Leu Phe Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn Gln
1670 1675 1680

Asp Ser Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu
1685 1690 1695

Thr Arg Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile
1700 1705 1710

Ala Leu Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr
1715 1720 1725

<210> 1549
<211> 2046
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1549
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

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Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320
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Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590
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Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Pro Gly Ala Ser Pro Gly
            740                 745                 750

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
        755                 760                 765

Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
    770                 775                 780

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser
785                 790                 795                 800

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
                805                 810                 815

Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
            820                 825                 830

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
        835                 840                 845

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
    850                 855                 860
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Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
865                 870                 875                 880

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser
                885                 890                 895

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
            900                 905                 910

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
        915                 920                 925

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly
    930                 935                 940

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
945                 950                 955                 960

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
                965                 970                 975

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly
            980                 985                 990

Thr Ser Ser Thr Gly Ser Pro Gly  Ser Ser Pro Ser Ala  Ser Thr Gly
        995                 1000                1005

Thr Gly  Pro Gly Thr Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro
    1010                1015                1020

Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Gly  Pro Pro Val
    1025                1030                1035

Leu Lys  Arg His Gln Arg Glu  Ile Thr Arg Thr Thr  Leu Gln Ser
    1040                1045                1050

Asp Gln  Glu Glu Ile Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met
    1055                1060                1065

Lys Lys  Glu Asp Phe Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser
    1070                1075                1080

Pro Arg  Ser Phe Gln Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala
    1085                1090                1095

Val Glu  Arg Leu Trp Asp Tyr  Gly Met Ser Ser Ser  Pro His Val
    1100                1105                1110

Leu Arg  Asn Arg Ala Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys
    1115                1120                1125
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Val Val  Phe Gln Glu Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu
    1130                1135                1140

Tyr Arg  Gly Glu Leu Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr
    1145                1150                1155

Ile Arg  Ala Glu Val Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn
    1160                1165                1170

Gln Ala  Ser Arg Pro Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr
    1175                1180                1185

Glu Glu  Asp Gln Arg Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val
    1190                1195                1200

Lys Pro  Asn Glu Thr Lys Thr  Tyr Phe Trp Lys Val  Gln His His
    1205                1210                1215

Met Ala  Pro Thr Lys Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr
    1220                1225                1230

Phe Ser  Asp Val Asp Leu Glu  Lys Asp Val His Ser  Gly Leu Ile
    1235                1240                1245

Gly Pro  Leu Leu Val Cys His  Thr Asn Thr Leu Asn  Pro Ala His
    1250                1255                1260

Gly Arg  Gln Val Thr Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile
    1265                1270                1275

Phe Asp  Glu Thr Lys Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg
    1280                1285                1290

Asn Cys  Arg Ala Pro Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe
    1295                1300                1305

Lys Glu  Asn Tyr Arg Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp
    1310                1315                1320

Thr Leu  Pro Gly Leu Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp
    1325                1330                1335

Tyr Leu  Leu Ser Met Gly Ser  Asn Glu Asn Ile His  Ser Ile His
    1340                1345                1350

Phe Ser  Gly His Val Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys
    1355                1360                1365

Met Ala  Leu Tyr Asn Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu
    1370                1375                1380
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Met Leu  Pro Ser Lys Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile
    1385                1390                1395

Gly Glu  His Leu His Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr
    1400                1405                1410

Ser Asn  Lys Cys Gln Thr Pro  Leu Gly Met Ala Ser  Gly His Ile
    1415                1420                1425

Arg Asp  Phe Gln Ile Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala
    1430                1435                1440

Pro Lys  Leu Ala Arg Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp
    1445                1450                1455

Ser Thr  Lys Glu Pro Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala
    1460                1465                1470

Pro Met  Ile Ile His Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys
    1475                1480                1485

Phe Ser  Ser Leu Tyr Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu
    1490                1495                1500

Asp Gly  Lys Lys Trp Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr
    1505                1510                1515

Leu Met  Val Phe Phe Gly Asn  Val Asp Ser Ser Gly  Ile Lys His
    1520                1525                1530

Asn Ile  Phe Asn Pro Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His
    1535                1540                1545

Pro Thr  His Tyr Ser Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met
    1550                1555                1560

Gly Cys  Asp Leu Asn Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser
    1565                1570                1575

Lys Ala  Ile Ser Asp Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr
    1580                1585                1590

Asn Met  Phe Ala Thr Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu
    1595                1600                1605

Gln Gly  Arg Ser Asn Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys
    1610                1615                1620

Glu Trp  Leu Gln Val Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly
    1625                1630                1635
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Val Thr  Thr Gln Gly Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val
    1640                1645                1650

Lys Glu  Phe Leu Ile Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr
    1655                1660                1665

Leu Phe  Phe Gln Asn Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln
    1670                1675                1680

Asp Ser  Phe Thr Pro Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu
    1685                1690                1695

Thr Arg  Tyr Leu Arg Ile His  Pro Gln Ser Trp Val  His Gln Ile
    1700                1705                1710

Ala Leu  Arg Met Glu Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr
    1715                1720                1725

Gly Ala  Gly Ser Pro Gly Ala  Glu Thr Ala Pro Gly  Ala Ser Pro
    1730                1735                1740

Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1745                1750                1755

Ser Thr  Gly Ser Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
    1760                1765                1770

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly
    1775                1780                1785

Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser Pro Gly  Ser Ser Thr
    1790                1795                1800

Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Thr Pro Gly  Ser Gly Thr
    1805                1810                1815

Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly
    1820                1825                1830

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly
    1835                1840                1845

Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly Pro Gly  Ser Ser Pro
    1850                1855                1860

Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ala Ser Pro  Gly Thr Ser
    1865                1870                1875

Ser Thr  Gly Ser Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
    1880                1885                1890
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Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly
    1895                 1900                 1905

Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly Pro Gly  Ser Ser Pro
    1910                 1915                 1920

Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ala Ser Pro  Gly Thr Ser
    1925                 1930                 1935

Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1940                 1945                 1950

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly
    1955                 1960                 1965

Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly Pro Gly  Ala Ser Pro
    1970                 1975                 1980

Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser
    1985                 1990                 1995

Thr Gly  Thr Gly Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
    2000                 2005                 2010

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Gly Ala
    2015                 2020                 2025

Glu Thr  Ala Glu Gln Lys Leu  Ile Ser Glu Glu Asp  Leu Ser Pro
    2030                 2035                 2040

Ala Thr  Gly
    2045


<210> 1550
<211> 2046
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
      polypeptide

<400> 1550
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1               5                  10                  15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro
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50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg

325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe

595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gly Pro Gly Ala Ser Pro Gly Thr Ser
740 745 750

Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
755 760 765

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser
770 775 780

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
785 790 795 800

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
805 810 815

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser
820 825 830

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly
835 840 845

Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
850 855 860

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser

865 870 875 880

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
885 890 895

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
900 905 910

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser
915 920 925

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser
930 935 940

Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
945 950 955 960

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
965 970 975

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser
980 985 990

Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
995 1000 1005

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser
1010 1015 1020

Ser Thr Pro Ser Gly Ala Thr Gly Ser Gly Gln Asn Pro Pro Val
1025 1030 1035

Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser
1040 1045 1050

Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
1055 1060 1065

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser
1070 1075 1080

Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala
1085 1090 1095

Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val
1100 1105 1110

Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys
1115 1120 1125

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu

1130 1135 1140

Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr
1145 1150 1155

Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn
1160 1165 1170

Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr
1175 1180 1185

Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val
1190 1195 1200

Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
1205 1210 1215

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr
1220 1225 1230

Phe Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile
1235 1240 1245

Gly Pro Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His
1250 1255 1260

Gly Arg Gln Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile
1265 1270 1275

Phe Asp Glu Thr Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg
1280 1285 1290

Asn Cys Arg Ala Pro Cys Asn Ile Gln Met Glu Asp Pro Thr Phe
1295 1300 1305

Lys Glu Asn Tyr Arg Phe His Ala Ile Asn Gly Tyr Ile Met Asp
1310 1315 1320

Thr Leu Pro Gly Leu Val Met Ala Gln Asp Gln Arg Ile Arg Trp
1325 1330 1335

Tyr Leu Leu Ser Met Gly Ser Asn Glu Asn Ile His Ser Ile His
1340 1345 1350

Phe Ser Gly His Val Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys
1355 1360 1365

Met Ala Leu Tyr Asn Leu Tyr Pro Gly Val Phe Glu Thr Val Glu
1370 1375 1380

Met Leu Pro Ser Lys Ala Gly Ile Trp Arg Val Glu Cys Leu Ile

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    1385                1390                1395

Gly Glu  His Leu His Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr
    1400                1405                1410

Ser Asn  Lys Cys Gln Thr Pro  Leu Gly Met Ala Ser  Gly His Ile
    1415                1420                1425

Arg Asp  Phe Gln Ile Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala
    1430                1435                1440

Pro Lys  Leu Ala Arg Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp
    1445                1450                1455

Ser Thr  Lys Glu Pro Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala
    1460                1465                1470

Pro Met  Ile Ile His Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys
    1475                1480                1485

Phe Ser  Ser Leu Tyr Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu
    1490                1495                1500

Asp Gly  Lys Lys Trp Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr
    1505                1510                1515

Leu Met  Val Phe Phe Gly Asn  Val Asp Ser Ser Gly  Ile Lys His
    1520                1525                1530

Asn Ile  Phe Asn Pro Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His
    1535                1540                1545

Pro Thr  His Tyr Ser Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met
    1550                1555                1560

Gly Cys  Asp Leu Asn Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser
    1565                1570                1575

Lys Ala  Ile Ser Asp Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr
    1580                1585                1590

Asn Met  Phe Ala Thr Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu
    1595                1600                1605

Gln Gly  Arg Ser Asn Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys
    1610                1615                1620

Glu Trp  Leu Gln Val Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly
    1625                1630                1635

Val Thr  Thr Gln Gly Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val
```

1640 1645 1650

Lys Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly His Gln Trp Thr
1655 1660 1665

Leu Phe Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn Gln
1670 1675 1680

Asp Ser Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu
1685 1690 1695

Thr Arg Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile
1700 1705 1710

Ala Leu Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr
1715 1720 1725

Gly Ala Gly Ser Pro Gly Ala Glu Thr Ala Pro Gly Ala Ser Pro
1730 1735 1740

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
1745 1750 1755

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
1760 1765 1770

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
1775 1780 1785

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
1790 1795 1800

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
1805 1810 1815

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
1820 1825 1830

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
1835 1840 1845

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
1850 1855 1860

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser
1865 1870 1875

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
1880 1885 1890

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly

1895 1900 1905

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
1910 1915 1920

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser
1925 1930 1935

Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
1940 1945 1950

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
1955 1960 1965

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
1970 1975 1980

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
1985 1990 1995

Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
2000 2005 2010

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Gly Ala
2015 2020 2025

Glu Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro
2030 2035 2040

Ala Thr Gly
2045

<210> 1551
<211> 2017
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1551
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

```
Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335
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Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605
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```
Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Gly Thr Ser Glu Ser Ala
            740                 745                 750

Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu
        755                 760                 765

Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
    770                 775                 780

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala
785                 790                 795                 800

Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
                805                 810                 815

Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr
            820                 825                 830

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr
        835                 840                 845

Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
    850                 855                 860

Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser
865                 870                 875                 880
```

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Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro
                885                 890                 895

Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
            900                 905                 910

Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr
        915                 920                 925

Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr
    930                 935                 940

Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu
945                 950                 955                 960

Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr
                965                 970                 975

Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro
            980                 985                 990

Ser Glu Gly Ser Ala Pro Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu
        995                 1000                1005

Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    1010                1015                1020

Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala Pro Gly  Pro Pro Val
    1025                1030                1035

Leu Lys  Arg His Gln Ala Glu  Ile Thr Arg Thr Thr  Leu Gln Ser
    1040                1045                1050

Asp Gln  Glu Glu Ile Asp Tyr  Asp Asp Thr Ile Ser  Val Glu Met
    1055                1060                1065

Lys Lys  Glu Asp Phe Asp Ile  Tyr Asp Glu Asp Glu  Asn Gln Ser
    1070                1075                1080

Pro Arg  Ser Phe Gln Lys Lys  Thr Arg His Tyr Phe  Ile Ala Ala
    1085                1090                1095

Val Glu  Arg Leu Trp Asp Tyr  Gly Met Ser Ser Ser  Pro His Val
    1100                1105                1110

Leu Arg  Asn Arg Ala Gln Ser  Gly Ser Val Pro Gln  Phe Lys Lys
    1115                1120                1125

Val Val  Phe Gln Glu Phe Thr  Asp Gly Ser Phe Thr  Gln Pro Leu
    1130                1135                1140
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```
Tyr Arg  Gly Glu Leu Asn Glu  His Leu Gly Leu Leu  Gly Pro Tyr
    1145                1150                1155

Ile Arg  Ala Glu Val Glu Asp  Asn Ile Met Val Thr  Phe Arg Asn
    1160                1165                1170

Gln Ala  Ser Arg Pro Tyr Ser  Phe Tyr Ser Ser Leu  Ile Ser Tyr
    1175                1180                1185

Glu Glu  Asp Gln Arg Gln Gly  Ala Glu Pro Arg Lys  Asn Phe Val
    1190                1195                1200

Lys Pro  Asn Glu Thr Lys Thr  Tyr Phe Trp Lys Val  Gln His His
    1205                1210                1215

Met Ala  Pro Thr Lys Asp Glu  Phe Asp Cys Lys Ala  Trp Ala Tyr
    1220                1225                1230

Phe Ser  Asp Val Asp Leu Glu  Lys Asp Val His Ser  Gly Leu Ile
    1235                1240                1245

Gly Pro  Leu Leu Val Cys His  Thr Asn Thr Leu Asn  Pro Ala His
    1250                1255                1260

Gly Arg  Gln Val Thr Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile
    1265                1270                1275

Phe Asp  Glu Thr Lys Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg
    1280                1285                1290

Asn Cys  Arg Ala Pro Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe
    1295                1300                1305

Lys Glu  Asn Tyr Arg Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp
    1310                1315                1320

Thr Leu  Pro Gly Leu Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp
    1325                1330                1335

Tyr Leu  Leu Ser Met Gly Ser  Asn Glu Asn Ile His  Ser Ile His
    1340                1345                1350

Phe Ser  Gly His Val Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys
    1355                1360                1365

Met Ala  Leu Tyr Asn Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu
    1370                1375                1380

Met Leu  Pro Ser Lys Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile
    1385                1390                1395
```

```
Gly Glu  His Leu His Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr
    1400                1405                1410

Ser Asn  Lys Cys Gln Thr Pro  Leu Gly Met Ala Ser  Gly His Ile
    1415                1420                1425

Arg Asp  Phe Gln Ile Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala
    1430                1435                1440

Pro Lys  Leu Ala Arg Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp
    1445                1450                1455

Ser Thr  Lys Glu Pro Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala
    1460                1465                1470

Pro Met  Ile Ile His Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys
    1475                1480                1485

Phe Ser  Ser Leu Tyr Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu
    1490                1495                1500

Asp Gly  Lys Lys Trp Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr
    1505                1510                1515

Leu Met  Val Phe Phe Gly Asn  Val Asp Ser Ser Gly  Ile Lys His
    1520                1525                1530

Asn Ile  Phe Asn Pro Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His
    1535                1540                1545

Pro Thr  His Tyr Ser Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met
    1550                1555                1560

Gly Cys  Asp Leu Asn Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser
    1565                1570                1575

Lys Ala  Ile Ser Asp Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr
    1580                1585                1590

Asn Met  Phe Ala Thr Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu
    1595                1600                1605

Gln Gly  Arg Ser Asn Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys
    1610                1615                1620

Glu Trp  Leu Gln Val Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly
    1625                1630                1635

Val Thr  Thr Gln Gly Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val
    1640                1645                1650
```

```
Lys Glu  Phe Leu Ile Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr
    1655                1660                1665

Leu Phe  Phe Gln Asn Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln
    1670                1675                1680

Asp Ser  Phe Thr Pro Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu
    1685                1690                1695

Thr Arg  Tyr Leu Arg Ile His  Pro Gln Ser Trp Val  His Gln Ile
    1700                1705                1710

Ala Leu  Arg Met Glu Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr
    1715                1720                1725

Gly Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1730                1735                1740

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser
    1745                1750                1755

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1760                1765                1770

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1775                1780                1785

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser
    1790                1795                1800

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser
    1805                1810                1815

Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly
    1820                1825                1830

Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1835                1840                1845

Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser
    1850                1855                1860

Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu Gly Thr  Ser Thr Glu
    1865                1870                1875

Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1880                1885                1890

Glu Ser  Gly Pro Gly Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly
    1895                1900                1905
```

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
1910 1915 1920

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala
1925 1930 1935

Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
1940 1945 1950

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
1955 1960 1965

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
1970 1975 1980

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
1985 1990 1995

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
2000 2005 2010

Gly Ser Ala Pro
2015

<210> 1552
<211> 2017
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1552
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala

100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His

370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr

645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gly Gly Thr Ser Glu Ser Ala Thr Pro
740 745 750

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
755 760 765

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
770 775 780

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
785 790 795 800

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
805 810 815

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
820 825 830

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
835 840 845

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
850 855 860

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala
865 870 875 880

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
885 890 895

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
900 905 910

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu

915 920 925

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
930 935 940

Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
945 950 955 960

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
965 970 975

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
980 985 990

Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
995 1000 1005

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
1010 1015 1020

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Gln Asn Pro Pro Val
1025 1030 1035

Leu Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln Ser
1040 1045 1050

Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
1055 1060 1065

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser
1070 1075 1080

Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala
1085 1090 1095

Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val
1100 1105 1110

Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys
1115 1120 1125

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu
1130 1135 1140

Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr
1145 1150 1155

Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn
1160 1165 1170

Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr

1175 1180 1185

Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val
1190 1195 1200

Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
1205 1210 1215

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr
1220 1225 1230

Phe Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile
1235 1240 1245

Gly Pro Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His
1250 1255 1260

Gly Arg Gln Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile
1265 1270 1275

Phe Asp Glu Thr Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg
1280 1285 1290

Asn Cys Arg Ala Pro Cys Asn Ile Gln Met Glu Asp Pro Thr Phe
1295 1300 1305

Lys Glu Asn Tyr Arg Phe His Ala Ile Asn Gly Tyr Ile Met Asp
1310 1315 1320

Thr Leu Pro Gly Leu Val Met Ala Gln Asp Gln Arg Ile Arg Trp
1325 1330 1335

Tyr Leu Leu Ser Met Gly Ser Asn Glu Asn Ile His Ser Ile His
1340 1345 1350

Phe Ser Gly His Val Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys
1355 1360 1365

Met Ala Leu Tyr Asn Leu Tyr Pro Gly Val Phe Glu Thr Val Glu
1370 1375 1380

Met Leu Pro Ser Lys Ala Gly Ile Trp Arg Val Glu Cys Leu Ile
1385 1390 1395

Gly Glu His Leu His Ala Gly Met Ser Thr Leu Phe Leu Val Tyr
1400 1405 1410

Ser Asn Lys Cys Gln Thr Pro Leu Gly Met Ala Ser Gly His Ile
1415 1420 1425

Arg Asp Phe Gln Ile Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala

1430 1435 1440

Pro Lys Leu Ala Arg Leu His Tyr Ser Gly Ser Ile Asn Ala Trp
1445 1450 1455

Ser Thr Lys Glu Pro Phe Ser Trp Ile Lys Val Asp Leu Leu Ala
1460 1465 1470

Pro Met Ile Ile His Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys
1475 1480 1485

Phe Ser Ser Leu Tyr Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu
1490 1495 1500

Asp Gly Lys Lys Trp Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr
1505 1510 1515

Leu Met Val Phe Phe Gly Asn Val Asp Ser Ser Gly Ile Lys His
1520 1525 1530

Asn Ile Phe Asn Pro Pro Ile Ile Ala Arg Tyr Ile Arg Leu His
1535 1540 1545

Pro Thr His Tyr Ser Ile Arg Ser Thr Leu Arg Met Glu Leu Met
1550 1555 1560

Gly Cys Asp Leu Asn Ser Cys Ser Met Pro Leu Gly Met Glu Ser
1565 1570 1575

Lys Ala Ile Ser Asp Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr
1580 1585 1590

Asn Met Phe Ala Thr Trp Ser Pro Ser Lys Ala Arg Leu His Leu
1595 1600 1605

Gln Gly Arg Ser Asn Ala Trp Arg Pro Gln Val Asn Asn Pro Lys
1610 1615 1620

Glu Trp Leu Gln Val Asp Phe Gln Lys Thr Met Lys Val Thr Gly
1625 1630 1635

Val Thr Thr Gln Gly Val Lys Ser Leu Leu Thr Ser Met Tyr Val
1640 1645 1650

Lys Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly His Gln Trp Thr
1655 1660 1665

Leu Phe Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn Gln
1670 1675 1680

Asp Ser Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu

1685 1690 1695

Thr Arg Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile
1700 1705 1710

Ala Leu Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr
1715 1720 1725

Gly Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
1730 1735 1740

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
1745 1750 1755

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
1760 1765 1770

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1775 1780 1785

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
1790 1795 1800

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
1805 1810 1815

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
1820 1825 1830

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1835 1840 1845

Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser
1850 1855 1860

Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu
1865 1870 1875

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
1880 1885 1890

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly
1895 1900 1905

Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser
1910 1915 1920

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala
1925 1930 1935

Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr

1940 1945 1950

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala
1955 1960 1965

Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser
1970 1975 1980

Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
1985 1990 1995

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
2000 2005 2010

Gly Ser Ala Pro
2015

<210> 1553
<211> 2045
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1553
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

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Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415
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Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685
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Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Pro Gly Ala Ser Pro Gly
            740                 745                 750

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
        755                 760                 765

Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
    770                 775                 780

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser
785                 790                 795                 800

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
                805                 810                 815

Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
            820                 825                 830

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
        835                 840                 845

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
    850                 855                 860

Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
865                 870                 875                 880

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser
                885                 890                 895

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
            900                 905                 910

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
        915                 920                 925

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly
    930                 935                 940

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
945                 950                 955                 960
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Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
                965                 970                 975

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly
            980                 985                 990

Thr Ser Ser Thr Gly Ser Pro Gly  Ser Ser Pro Ser Ala  Ser Thr Gly
        995                 1000                1005

Thr Gly  Pro Gly Thr Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro
    1010                1015                1020

Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Pro Val Leu
    1025                1030                1035

Lys Arg  His Gln Ala Glu Ile  Thr Arg Thr Thr Leu  Gln Ser Asp
    1040                1045                1050

Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile Ser Val  Glu Met Lys
    1055                1060                1065

Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp Glu Asn  Gln Ser Pro
    1070                1075                1080

Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr Phe Ile  Ala Ala Val
    1085                1090                1095

Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser Ser Pro  His Val Leu
    1100                1105                1110

Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro Gln Phe  Lys Lys Val
    1115                1120                1125

Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe Thr Gln  Pro Leu Tyr
    1130                1135                1140

Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu Leu Gly  Pro Tyr Ile
    1145                1150                1155

Arg Ala  Glu Val Glu Asp Asn  Ile Met Val Thr Phe  Arg Asn Gln
    1160                1165                1170

Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser Leu Ile  Ser Tyr Glu
    1175                1180                1185

Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg Lys Asn  Phe Val Lys
    1190                1195                1200

Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys Val Gln  His His Met
    1205                1210                1215
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Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys Ala Trp  Ala Tyr Phe
    1220                1225                1230

Ser Asp  Val Asp Leu Glu Lys  Asp Val His Ser Gly  Leu Ile Gly
    1235                1240                1245

Pro Leu  Leu Val Cys His Thr  Asn Thr Leu Asn Pro  Ala His Gly
    1250                1255                1260

Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu Phe Phe  Thr Ile Phe
    1265                1270                1275

Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu Asn Met  Glu Arg Asn
    1280                1285                1290

Cys Arg  Ala Pro Cys Asn Ile  Gln Met Glu Asp Pro  Thr Phe Lys
    1295                1300                1305

Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly Tyr Ile  Met Asp Thr
    1310                1315                1320

Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln Arg Ile  Arg Trp Tyr
    1325                1330                1335

Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile His Ser  Ile His Phe
    1340                1345                1350

Ser Gly  His Val Phe Thr Val  Arg Lys Lys Glu Glu  Tyr Lys Met
    1355                1360                1365

Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe Glu Thr  Val Glu Met
    1370                1375                1380

Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val Glu Cys  Leu Ile Gly
    1385                1390                1395

Glu His  Leu His Ala Gly Met  Ser Thr Leu Phe Leu  Val Tyr Ser
    1400                1405                1410

Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala Ser Gly  His Ile Arg
    1415                1420                1425

Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr Gly Gln  Trp Ala Pro
    1430                1435                1440

Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser Ile Asn  Ala Trp Ser
    1445                1450                1455

Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val Asp Leu  Leu Ala Pro
    1460                1465                1470
```

```
Met Ile  Ile His Gly Ile Lys  Thr Gln Gly Ala Arg  Gln Lys Phe
    1475                 1480                 1485

Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile Met Tyr  Ser Leu Asp
    1490                 1495                 1500

Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn Ser Thr  Gly Thr Leu
    1505                 1510                 1515

Met Val  Phe Phe Gly Asn Val  Asp Ser Ser Gly Ile  Lys His Asn
    1520                 1525                 1530

Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr Ile Arg  Leu His Pro
    1535                 1540                 1545

Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg Met Glu  Leu Met Gly
    1550                 1555                 1560

Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu Gly Met  Glu Ser Lys
    1565                 1570                 1575

Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser Ser Tyr  Phe Thr Asn
    1580                 1585                 1590

Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala Arg Leu  His Leu Gln
    1595                 1600                 1605

Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val Asn Asn  Pro Lys Glu
    1610                 1615                 1620

Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met Lys Val  Thr Gly Val
    1625                 1630                 1635

Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr Ser Met  Tyr Val Lys
    1640                 1645                 1650

Glu Phe  Leu Ile Ser Ser Ser  Gln Asp Gly His Gln  Trp Thr Leu
    1655                 1660                 1665

Phe Phe  Gln Asn Gly Lys Val  Lys Val Phe Gln Gly  Asn Gln Asp
    1670                 1675                 1680

Ser Phe  Thr Pro Val Val Asn  Ser Leu Asp Pro Pro  Leu Leu Thr
    1685                 1690                 1695

Arg Tyr  Leu Arg Ile His Pro  Gln Ser Trp Val His  Gln Ile Ala
    1700                 1705                 1710

Leu Arg  Met Glu Val Leu Gly  Cys Glu Ala Gln Asp  Leu Tyr Gly
    1715                 1720                 1725
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```
Ala Gly  Ser Pro Gly Ala Glu  Thr Ala Pro Gly Ala  Ser Pro Gly
    1730                1735                1740

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1745                1750                1755

Thr Gly  Ser Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1760                1765                1770

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Thr
    1775                1780                1785

Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro
    1790                1795                1800

Ser Gly  Ala Thr Gly Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala
    1805                1810                1815

Ser Ser  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1820                1825                1830

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser
    1835                1840                1845

Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro Gly Ser  Ser Pro Ser
    1850                1855                1860

Ala Ser  Thr Gly Thr Gly Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1865                1870                1875

Thr Gly  Ser Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1880                1885                1890

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser
    1895                1900                1905

Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro Gly Ser  Ser Pro Ser
    1910                1915                1920

Ala Ser  Thr Gly Thr Gly Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1925                1930                1935

Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    1940                1945                1950

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser
    1955                1960                1965

Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro Gly Ala  Ser Pro Gly
    1970                1975                1980
```

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
1985 1990 1995

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
2000 2005 2010

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Gly Ala Glu
2015 2020 2025

Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala
2030 2035 2040

Thr Gly
2045

<210> 1554
<211> 2045
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1554
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser

145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr

420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala

690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gly Pro Gly Ala Ser Pro Gly Thr Ser
740 745 750

Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
755 760 765

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser
770 775 780

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
785 790 795 800

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
805 810 815

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser
820 825 830

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly
835 840 845

Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
850 855 860

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
865 870 875 880

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
885 890 895

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
900 905 910

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser
915 920 925

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser
930 935 940

Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
945 950 955 960

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser

965 970 975

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser
980 985 990

Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
995 1000 1005

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser
1010 1015 1020

Ser Thr Pro Ser Gly Ala Thr Gly Ser Gln Asn Pro Pro Val Leu
1025 1030 1035

Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp
1040 1045 1050

Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys
1055 1060 1065

Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
1070 1075 1080

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val
1085 1090 1095

Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu
1100 1105 1110

Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val
1115 1120 1125

Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
1130 1135 1140

Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile
1145 1150 1155

Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln
1160 1165 1170

Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu
1175 1180 1185

Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys
1190 1195 1200

Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His Met
1205 1210 1215

Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe

1220 1225 1230

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly
1235 1240 1245

Pro Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly
1250 1255 1260

Arg Gln Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe
1265 1270 1275

Asp Glu Thr Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn
1280 1285 1290

Cys Arg Ala Pro Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys
1295 1300 1305

Glu Asn Tyr Arg Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr
1310 1315 1320

Leu Pro Gly Leu Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr
1325 1330 1335

Leu Leu Ser Met Gly Ser Asn Glu Asn Ile His Ser Ile His Phe
1340 1345 1350

Ser Gly His Val Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met
1355 1360 1365

Ala Leu Tyr Asn Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met
1370 1375 1380

Leu Pro Ser Lys Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly
1385 1390 1395

Glu His Leu His Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser
1400 1405 1410

Asn Lys Cys Gln Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg
1415 1420 1425

Asp Phe Gln Ile Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro
1430 1435 1440

Lys Leu Ala Arg Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser
1445 1450 1455

Thr Lys Glu Pro Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro
1460 1465 1470

Met Ile Ile His Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe

1475 1480 1485

Ser Ser Leu Tyr Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp
1490 1495 1500

Gly Lys Lys Trp Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu
1505 1510 1515

Met Val Phe Phe Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn
1520 1525 1530

Ile Phe Asn Pro Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro
1535 1540 1545

Thr His Tyr Ser Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly
1550 1555 1560

Cys Asp Leu Asn Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys
1565 1570 1575

Ala Ile Ser Asp Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn
1580 1585 1590

Met Phe Ala Thr Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln
1595 1600 1605

Gly Arg Ser Asn Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu
1610 1615 1620

Trp Leu Gln Val Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val
1625 1630 1635

Thr Thr Gln Gly Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys
1640 1645 1650

Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu
1655 1660 1665

Phe Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp
1670 1675 1680

Ser Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr
1685 1690 1695

Arg Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile Ala
1700 1705 1710

Leu Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly
1715 1720 1725

Ala Gly Ser Pro Gly Ala Glu Thr Ala Pro Gly Ala Ser Pro Gly

1730 1735 1740

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
1745 1750 1755

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
1760 1765 1770

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr
1775 1780 1785

Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
1790 1795 1800

Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
1805 1810 1815

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
1820 1825 1830

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser
1835 1840 1845

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser
1850 1855 1860

Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
1865 1870 1875

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
1880 1885 1890

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser
1895 1900 1905

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser
1910 1915 1920

Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
1925 1930 1935

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
1940 1945 1950

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser
1955 1960 1965

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly
1970 1975 1980

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr

1985 1990 1995

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
2000 2005 2010

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Gly Ala Glu
2015 2020 2025

Thr Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Ser Pro Ala
2030 2035 2040

Thr Gly
2045

<210> 1555
<211> 2301
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1555
Met Gln Ile Glu Leu Ser Thr Cys Phe Phe Leu Cys Leu Leu Arg Phe
1 5 10 15

Cys Phe Ser Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
20 25 30

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
35 40 45

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
50 55 60

Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
65 70 75 80

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
85 90 95

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
100 105 110

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
115 120 125

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly
130 135 140

Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser
145 150 155 160

```
Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
                165                 170                 175

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser
            180                 185                 190

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
        195                 200                 205

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
    210                 215                 220

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly
225                 230                 235                 240

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
                245                 250                 255

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            260                 265                 270

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
        275                 280                 285

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    290                 295                 300

Ser Ala Pro Gln Ser Pro Arg Ser Phe Gln Gly Pro Glu Gly Pro Ser
305                 310                 315                 320

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
                325                 330                 335

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            340                 345                 350

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        355                 360                 365

Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro
    370                 375                 380

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
385                 390                 395                 400

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                405                 410                 415

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            420                 425                 430
```

```
Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        435                 440                 445

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    450                 455                 460

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
465                 470                 475                 480

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                485                 490                 495

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            500                 505                 510

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        515                 520                 525

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    530                 535                 540

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
545                 550                 555                 560

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                565                 570                 575

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            580                 585                 590

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        595                 600                 605

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    610                 615                 620

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
625                 630                 635                 640

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                645                 650                 655

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            660                 665                 670

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        675                 680                 685

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    690                 695                 700
```

```
Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
705                 710                 715                 720

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                725                 730                 735

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            740                 745                 750

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        755                 760                 765

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    770                 775                 780

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
785                 790                 795                 800

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                805                 810                 815

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            820                 825                 830

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        835                 840                 845

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    850                 855                 860

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
865                 870                 875                 880

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                885                 890                 895

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            900                 905                 910

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        915                 920                 925

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    930                 935                 940

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
945                 950                 955                 960

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                965                 970                 975
```

```
Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            980                 985                 990

Phe Ser Gly Glu Thr Val Phe Met  Ser Met Glu Asn Pro  Gly Leu Trp
        995                 1000                1005

Ile Leu  Gly Cys His Asn Ser  Asp Phe Arg Asn Arg  Gly Met Thr
    1010                1015                1020

Ala Leu  Leu Lys Val Ser Ser  Cys Asp Lys Asn Thr  Gly Asp Tyr
    1025                1030                1035

Tyr Glu  Asp Ser Tyr Glu Asp  Ile Ser Ala Tyr Leu  Leu Ser Lys
    1040                1045                1050

Asn Asn  Ala Ile Glu Pro Arg  Ser Phe Ser Gln Asn  Gly Thr Ser
    1055                1060                1065

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
    1070                1075                1080

Ser Gly  Ser Glu Thr Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1085                1090                1095

Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro
    1100                1105                1110

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1115                1120                1125

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1130                1135                1140

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1145                1150                1155

Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro
    1160                1165                1170

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Pro
    1175                1180                1185

Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Ser Pro  Ala Gly Ser
    1190                1195                1200

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1205                1210                1215

Ser Ala  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1220                1225                1230
```

```
Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1235                 1240                 1245

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
    1250                 1255                 1260

Ser Gly  Ser Glu Thr Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1265                 1270                 1275

Glu Thr  Pro Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1280                 1285                 1290

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1295                 1300                 1305

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Glu  Pro Ala Thr
    1310                 1315                 1320

Ser Gly  Ser Glu Thr Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1325                 1330                 1335

Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1340                 1345                 1350

Gln Ser  Pro Arg Ser Phe Gln  Lys Lys Thr Arg His  Tyr Phe Ile
    1355                 1360                 1365

Ala Ala  Val Glu Arg Leu Trp  Asp Tyr Gly Met Ser  Ser Ser Pro
    1370                 1375                 1380

His Val  Leu Arg Asn Arg Ala  Gln Ser Gly Ser Val  Pro Gln Phe
    1385                 1390                 1395

Lys Lys  Val Val Phe Gln Glu  Phe Thr Asp Gly Ser  Phe Thr Gln
    1400                 1405                 1410

Pro Leu  Tyr Arg Gly Glu Leu  Asn Glu His Leu Gly  Leu Leu Gly
    1415                 1420                 1425

Pro Tyr  Ile Arg Ala Glu Val  Glu Asp Asn Ile Met  Val Thr Phe
    1430                 1435                 1440

Arg Asn  Gln Ala Ser Arg Pro  Tyr Ser Phe Tyr Ser  Ser Leu Ile
    1445                 1450                 1455

Ser Tyr  Glu Glu Asp Gln Arg  Gln Gly Ala Glu Pro  Arg Lys Asn
    1460                 1465                 1470

Phe Val  Lys Pro Asn Glu Thr  Lys Thr Tyr Phe Trp  Lys Val Gln
    1475                 1480                 1485
```

```
His His  Met Ala Pro Thr Lys  Asp Glu Phe Asp Cys  Lys Ala Trp
    1490                1495                1500

Ala Tyr  Phe Ser Asp Val Asp  Leu Glu Lys Asp Val  His Ser Gly
    1505                1510                1515

Leu Ile  Gly Pro Leu Leu Val  Cys His Thr Asn Thr  Leu Asn Pro
    1520                1525                1530

Ala His  Gly Arg Gln Val Thr  Val Gln Glu Phe Ala  Leu Phe Phe
    1535                1540                1545

Thr Ile  Phe Asp Glu Thr Lys  Ser Trp Tyr Phe Thr  Glu Asn Met
    1550                1555                1560

Glu Arg  Asn Cys Arg Ala Pro  Cys Asn Ile Gln Met  Glu Asp Pro
    1565                1570                1575

Thr Phe  Lys Glu Asn Tyr Arg  Phe His Ala Ile Asn  Gly Tyr Ile
    1580                1585                1590

Met Asp  Thr Leu Pro Gly Leu  Val Met Ala Gln Asp  Gln Arg Ile
    1595                1600                1605

Arg Trp  Tyr Leu Leu Ser Met  Gly Ser Asn Glu Asn  Ile His Ser
    1610                1615                1620

Ile His  Phe Ser Gly His Val  Phe Thr Val Arg Lys  Lys Glu Glu
    1625                1630                1635

Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro Gly Val  Phe Glu Thr
    1640                1645                1650

Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile Trp Arg  Val Glu Cys
    1655                1660                1665

Leu Ile  Gly Glu His Leu His  Ala Gly Met Ser Thr  Leu Phe Leu
    1670                1675                1680

Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu Gly Met  Ala Ser Gly
    1685                1690                1695

His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser Gly Gln  Tyr Gly Gln
    1700                1705                1710

Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr Ser Gly  Ser Ile Asn
    1715                1720                1725

Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp Ile Lys  Val Asp Leu
    1730                1735                1740
```

```
Leu Ala  Pro Met Ile Ile His  Gly Ile Lys Thr Gln  Gly Ala Arg
    1745                1750                1755

Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln Phe Ile  Ile Met Tyr
    1760                1765                1770

Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr Arg Gly  Asn Ser Thr
    1775                1780                1785

Gly Thr  Leu Met Val Phe Phe  Gly Asn Val Asp Ser  Ser Gly Ile
    1790                1795                1800

Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile Ala Arg  Tyr Ile Arg
    1805                1810                1815

Leu His  Pro Thr His Tyr Ser  Ile Arg Ser Thr Leu  Arg Met Glu
    1820                1825                1830

Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser Met Pro  Leu Gly Met
    1835                1840                1845

Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile Thr Ala  Ser Ser Tyr
    1850                1855                1860

Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro Ser Lys  Ala Arg Leu
    1865                1870                1875

His Leu  Gln Gly Arg Ser Asn  Ala Trp Arg Pro Gln  Val Asn Asn
    1880                1885                1890

Pro Lys  Glu Trp Leu Gln Val  Asp Phe Gln Lys Thr  Met Lys Val
    1895                1900                1905

Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser Leu Leu  Thr Ser Met
    1910                1915                1920

Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser Gln Asp  Gly His Gln
    1925                1930                1935

Trp Thr  Leu Phe Phe Gln Asn  Gly Lys Val Lys Val  Phe Gln Gly
    1940                1945                1950

Asn Gln  Asp Ser Phe Thr Pro  Val Val Asn Ser Leu  Asp Pro Pro
    1955                1960                1965

Leu Leu  Thr Arg Tyr Leu Arg  Ile His Pro Gln Ser  Trp Val His
    1970                1975                1980

Gln Ile  Ala Leu Arg Met Glu  Val Leu Gly Cys Glu  Ala Gln Asp
    1985                1990                1995
```

```
Leu Tyr  Gly Pro Glu Gly Pro  Ser Gln Ser Pro Arg  Ser Phe Gln
    2000                 2005                 2010

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    2015                 2020                 2025

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    2030                 2035                 2040

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    2045                 2050                 2055

Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    2060                 2065                 2070

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Ser Pro
    2075                 2080                 2085

Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Thr Ser  Glu Ser Ala
    2090                 2095                 2100

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    2105                 2110                 2115

Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    2120                 2125                 2130

Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr Glu Glu  Gly Ser Pro
    2135                 2140                 2145

Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Thr Ser  Thr Glu Pro
    2150                 2155                 2160

Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    2165                 2170                 2175

Ser Gly  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    2180                 2185                 2190

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    2195                 2200                 2205

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Ser Glu  Pro Ala Thr
    2210                 2215                 2220

Ser Gly  Ser Glu Thr Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    2225                 2230                 2235

Thr Glu  Glu Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    2240                 2245                 2250
```

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu
2255 2260 2265

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala
2270 2275 2280

Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
2285 2290 2295

Ser Ala Pro
2300

<210> 1556
<211> 2876
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1556
Met Gln Ile Glu Leu Ser Thr Cys Phe Phe Leu Cys Leu Leu Arg Phe
1 5 10 15

Cys Phe Ser Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
20 25 30

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
35 40 45

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
50 55 60

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
65 70 75 80

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
85 90 95

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
100 105 110

Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
115 120 125

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
130 135 140

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
145 150 155 160

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly

```
                165                 170                 175

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu
            180                 185                 190

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
        195                 200                 205

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
    210                 215                 220

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
225                 230                 235                 240

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
                245                 250                 255

Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            260                 265                 270

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
        275                 280                 285

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
    290                 295                 300

Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
305                 310                 315                 320

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
                325                 330                 335

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
            340                 345                 350

Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
        355                 360                 365

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
    370                 375                 380

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
385                 390                 395                 400

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
                405                 410                 415

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
            420                 425                 430

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
```

435 440 445

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
450 455 460

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
465 470 475 480

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
485 490 495

Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
500 505 510

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
515 520 525

Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
530 535 540

Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
545 550 555 560

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
565 570 575

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
580 585 590

Ser Ala Pro Gln Ser Pro Arg Ser Phe Gln Gly Pro Ser Gly Pro Ala
595 600 605

Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr Met
610 615 620

Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro Arg
625 630 635 640

Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys Thr
645 650 655

Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro Arg
660 665 670

Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val Tyr
675 680 685

Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val Ser
690 695 700

Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala Glu

```
705                 710                 715                 720

Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val Phe
                725                 730                 735

Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn Gly
            740                 745                 750

Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser His
        755                 760                 765

Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu Leu
    770                 775                 780

Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu His
785                 790                 795                 800

Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp His
                805                 810                 815

Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser Ala
            820                 825                 830

Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg Ser
        835                 840                 845

Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His Val
    850                 855                 860

Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu Gly
865                 870                 875                 880

His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile Ser
                885                 890                 895

Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly Gln
            900                 905                 910

Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met Glu
        915                 920                 925

Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg Met
    930                 935                 940

Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp Ser
945                 950                 955                 960

Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe Ile
                965                 970                 975

Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His Tyr
```

980 985 990

Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu Ala
995 1000 1005

Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
1010 1015 1020

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr
1025 1030 1035

Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser
1040 1045 1050

Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu
1055 1060 1065

Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr
1070 1075 1080

Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu
1085 1090 1095

Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly
1100 1105 1110

Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro
1115 1120 1125

Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe
1130 1135 1140

Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu
1145 1150 1155

Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile
1160 1165 1170

Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu
1175 1180 1185

Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro
1190 1195 1200

Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser
1205 1210 1215

Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln
1220 1225 1230

Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser

1235 1240 1245

Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
1250 1255 1260

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe
1265 1270 1275

Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly
1280 1285 1290

Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly
1295 1300 1305

Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly
1310 1315 1320

Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu
1325 1330 1335

Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly
1340 1345 1350

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
1355 1360 1365

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser
1370 1375 1380

Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr
1385 1390 1395

Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
1400 1405 1410

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
1415 1420 1425

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
1430 1435 1440

Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu
1445 1450 1455

Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
1460 1465 1470

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
1475 1480 1485

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser

1490 1495 1500

Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr
1505 1510 1515

Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
1520 1525 1530

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
1535 1540 1545

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser
1550 1555 1560

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
1565 1570 1575

Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
1580 1585 1590

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
1595 1600 1605

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
1610 1615 1620

Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
1625 1630 1635

Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
1640 1645 1650

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
1655 1660 1665

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
1670 1675 1680

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
1685 1690 1695

Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
1700 1705 1710

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
1715 1720 1725

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
1730 1735 1740

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser

1745 1750 1755

Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
1760 1765 1770

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
1775 1780 1785

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
1790 1795 1800

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
1805 1810 1815

Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
1820 1825 1830

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
1835 1840 1845

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
1850 1855 1860

Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr
1865 1870 1875

Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
1880 1885 1890

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
1895 1900 1905

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser
1910 1915 1920

Glu Gly Ser Ala Pro Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr
1925 1930 1935

Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
1940 1945 1950

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly
1955 1960 1965

Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp
1970 1975 1980

Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His
1985 1990 1995

Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn

2000 2005 2010

Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe
2015 2020 2025

Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala
2030 2035 2040

Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr Lys Thr Tyr
2045 2050 2055

Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp Glu Phe
2060 2065 2070

Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu Lys
2075 2080 2085

Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val Cys His Thr
2090 2095 2100

Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr Val Gln Glu
2105 2110 2115

Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys Ser Trp Tyr
2120 2125 2130

Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro Cys Asn Ile
2135 2140 2145

Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg Phe His Ala
2150 2155 2160

Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu Val Met Ala
2165 2170 2175

Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met Gly Ser Asn
2180 2185 2190

Glu Asn Ile His Ser Ile His Phe Ser Gly His Val Phe Thr Val
2195 2200 2205

Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn Leu Tyr Pro
2210 2215 2220

Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys Ala Gly Ile
2225 2230 2235

Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His Ala Gly Met
2240 2245 2250

Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln Thr Pro Leu

2255 2260 2265

Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile Thr Ala Ser
2270 2275 2280

Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg Leu His Tyr
2285 2290 2295

Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro Phe Ser Trp
2300 2305 2310

Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His Gly Ile Lys
2315 2320 2325

Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr Ile Ser Gln
2330 2335 2340

Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp Gln Thr Tyr
2345 2350 2355

Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe Gly Asn Val
2360 2365 2370

Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro Pro Ile Ile
2375 2380 2385

Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser Ile Arg Ser
2390 2395 2400

Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn Ser Cys Ser
2405 2410 2415

Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp Ala Gln Ile
2420 2425 2430

Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp Ser Pro
2435 2440 2445

Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg
2450 2455 2460

Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln
2465 2470 2475

Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser
2480 2485 2490

Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser
2495 2500 2505

Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val

2510 2515 2520

Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn
2525 2530 2535

Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro
2540 2545 2550

Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly
2555 2560 2565

Cys Glu Ala Gln Asp Leu Tyr Gly Pro Glu Gly Pro Ser Gln Ser
2570 2575 2580

Pro Arg Ser Phe Gln Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
2585 2590 2595

Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
2600 2605 2610

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
2615 2620 2625

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
2630 2635 2640

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
2645 2650 2655

Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
2660 2665 2670

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
2675 2680 2685

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
2690 2695 2700

Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr
2705 2710 2715

Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
2720 2725 2730

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
2735 2740 2745

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
2750 2755 2760

Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser

2765 2770 2775

Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
2780 2785 2790

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
2795 2800 2805

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
2810 2815 2820

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
2825 2830 2835

Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
2840 2845 2850

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
2855 2860 2865

Glu Pro Ser Glu Gly Ser Ala Pro
2870 2875

<210> 1557
<211> 2576
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1557
Met Gln Ile Glu Leu Ser Thr Cys Phe Phe Leu Cys Leu Leu Arg Phe
1 5 10 15

Cys Phe Ser Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser
20 25 30

Trp Asp Tyr Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg
35 40 45

Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val
50 55 60

Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile
65 70 75 80

Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln
85 90 95

Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser
100 105 110

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His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser
        115                 120                 125

Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp
    130                 135                 140

Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu
145                 150                 155                 160

Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser
                165                 170                 175

Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile
            180                 185                 190

Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr
        195                 200                 205

Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly
    210                 215                 220

Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp
225                 230                 235                 240

Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr
                245                 250                 255

Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val
            260                 265                 270

Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile
        275                 280                 285

Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser
    290                 295                 300

Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met
305                 310                 315                 320

Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His
                325                 330                 335

Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro
            340                 345                 350

Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp
        355                 360                 365

Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser
    370                 375                 380
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Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr
385                 390                 395                 400

Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro
                405                 410                 415

Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn
            420                 425                 430

Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met
        435                 440                 445

Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu
    450                 455                 460

Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu
465                 470                 475                 480

Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro
                485                 490                 495

His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys
            500                 505                 510

Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe
        515                 520                 525

Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp
    530                 535                 540

Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg
545                 550                 555                 560

Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu
                565                 570                 575

Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val
            580                 585                 590

Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu
        595                 600                 605

Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp
    610                 615                 620

Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val
625                 630                 635                 640

Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp
                645                 650                 655
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Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe
            660                 665                 670

Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr
        675                 680                 685

Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro
    690                 695                 700

Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly
705                 710                 715                 720

Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp
                725                 730                 735

Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys
            740                 745                 750

Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Ser Pro Ala
        755                 760                 765

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
    770                 775                 780

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
785                 790                 795                 800

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr
                805                 810                 815

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
            820                 825                 830

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        835                 840                 845

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
    850                 855                 860

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
865                 870                 875                 880

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
                885                 890                 895

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
            900                 905                 910

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
        915                 920                 925
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Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    930                 935                 940

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
945                 950                 955                 960

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
                965                 970                 975

Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
            980                 985                 990

Gly Ser Glu Pro Ala Thr Ser Gly  Ser Glu Thr Pro Gly  Thr Ser Thr
        995                 1000                1005

Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser
    1010                1015                1020

Glu Gly  Ser Ala Pro Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1025                1030                1035

Gly Pro  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    1040                1045                1050

Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Glu
    1055                1060                1065

Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser
    1070                1075                1080

Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1085                1090                1095

Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    1100                1105                1110

Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr
    1115                1120                1125

Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser
    1130                1135                1140

Glu Gly  Ser Ala Pro Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    1145                1150                1155

Ala Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    1160                1165                1170

Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr
    1175                1180                1185
```

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Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr Ser Glu  Ser Ala Thr
    1190                 1195                 1200

Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu
    1205                 1210                 1215

Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    1220                 1225                 1230

Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr Pro Gly  Thr Ser Glu
    1235                 1240                 1245

Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser
    1250                 1255                 1260

Glu Gly  Ser Ala Pro Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1265                 1270                 1275

Gly Pro  Gly Ser Pro Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly
    1280                 1285                 1290

Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala
    1295                 1300                 1305

Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr Ser Glu  Ser Ala Thr
    1310                 1315                 1320

Pro Glu  Ser Gly Pro Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    1325                 1330                 1335

Ala Pro  Gln Ser Pro Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr
    1340                 1345                 1350

Phe Ile  Ala Ala Val Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser
    1355                 1360                 1365

Ser Pro  His Val Leu Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro
    1370                 1375                 1380

Gln Phe  Lys Lys Val Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe
    1385                 1390                 1395

Thr Gln  Pro Leu Tyr Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu
    1400                 1405                 1410

Leu Gly  Pro Tyr Ile Arg Ala  Glu Val Glu Asp Asn  Ile Met Val
    1415                 1420                 1425

Thr Phe  Arg Asn Gln Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser
    1430                 1435                 1440
```

```
Leu Ile  Ser Tyr Glu Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg
    1445                 1450                 1455

Lys Asn  Phe Val Lys Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys
    1460                 1465                 1470

Val Gln  His His Met Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys
    1475                 1480                 1485

Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu Glu Lys  Asp Val His
    1490                 1495                 1500

Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys His Thr  Asn Thr Leu
    1505                 1510                 1515

Asn Pro  Ala His Gly Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu
    1520                 1525                 1530

Phe Phe  Thr Ile Phe Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu
    1535                 1540                 1545

Asn Met  Glu Arg Asn Cys Arg  Ala Pro Cys Asn Ile  Gln Met Glu
    1550                 1555                 1560

Asp Pro  Thr Phe Lys Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly
    1565                 1570                 1575

Tyr Ile  Met Asp Thr Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln
    1580                 1585                 1590

Arg Ile  Arg Trp Tyr Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile
    1595                 1600                 1605

His Ser  Ile His Phe Ser Gly  His Val Phe Thr Val  Arg Lys Lys
    1610                 1615                 1620

Glu Glu  Tyr Lys Met Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe
    1625                 1630                 1635

Glu Thr  Val Glu Met Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val
    1640                 1645                 1650

Glu Cys  Leu Ile Gly Glu His  Leu His Ala Gly Met  Ser Thr Leu
    1655                 1660                 1665

Phe Leu  Val Tyr Ser Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala
    1670                 1675                 1680

Ser Gly  His Ile Arg Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr
    1685                 1690                 1695
```

```
Gly Gln  Trp Ala Pro Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser
    1700                1705                1710

Ile Asn  Ala Trp Ser Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val
    1715                1720                1725

Asp Leu  Leu Ala Pro Met Ile  Ile His Gly Ile Lys  Thr Gln Gly
    1730                1735                1740

Ala Arg  Gln Lys Phe Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile
    1745                1750                1755

Met Tyr  Ser Leu Asp Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn
    1760                1765                1770

Ser Thr  Gly Thr Leu Met Val  Phe Phe Gly Asn Val  Asp Ser Ser
    1775                1780                1785

Gly Ile  Lys His Asn Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr
    1790                1795                1800

Ile Arg  Leu His Pro Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg
    1805                1810                1815

Met Glu  Leu Met Gly Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu
    1820                1825                1830

Gly Met  Glu Ser Lys Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser
    1835                1840                1845

Ser Tyr  Phe Thr Asn Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala
    1850                1855                1860

Arg Leu  His Leu Gln Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val
    1865                1870                1875

Asn Asn  Pro Lys Glu Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met
    1880                1885                1890

Lys Val  Thr Gly Val Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr
    1895                1900                1905

Ser Met  Tyr Val Lys Glu Phe  Leu Ile Ser Ser Ser  Gln Asp Gly
    1910                1915                1920

His Gln  Trp Thr Leu Phe Phe  Gln Asn Gly Lys Val  Lys Val Phe
    1925                1930                1935

Gln Gly  Asn Gln Asp Ser Phe  Thr Pro Val Val Asn  Ser Leu Asp
    1940                1945                1950
```

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Pro Pro  Leu Leu Thr Arg Tyr  Leu Arg Ile His Pro  Gln Ser Trp
    1955                 1960                 1965

Val His  Gln Ile Ala Leu Arg  Met Glu Val Leu Gly  Cys Glu Ala
    1970                 1975                 1980

Gln Asp  Leu Tyr Gly Pro Glu  Gly Pro Ser Gln Ser  Pro Arg Ser
    1985                 1990                 1995

Phe Gln  Gly Ser Pro Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly
    2000                 2005                 2010

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Thr
    2015                 2020                 2025

Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser Pro Ala  Gly Ser Pro
    2030                 2035                 2040

Thr Ser  Thr Glu Glu Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    2045                 2050                 2055

Ala Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    2060                 2065                 2070

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro
    2075                 2080                 2085

Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Ser Glu Pro  Ala Thr Ser
    2090                 2095                 2100

Gly Ser  Glu Thr Pro Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr
    2105                 2110                 2115

Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    2120                 2125                 2130

Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr
    2135                 2140                 2145

Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser Pro Ala  Gly Ser Pro
    2150                 2155                 2160

Thr Ser  Thr Glu Glu Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    2165                 2170                 2175

Ala Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    2180                 2185                 2190

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Thr
    2195                 2200                 2205
```

```
Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr Ser Glu  Ser Ala Thr
    2210                 2215                 2220

Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu
    2225                 2230                 2235

Thr Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    2240                 2245                 2250

Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Glu
    2255                 2260                 2265

Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr Ser Glu  Ser Ala Thr
    2270                 2275                 2280

Pro Glu  Ser Gly Pro Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr
    2285                 2290                 2295

Glu Glu  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    2300                 2305                 2310

Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr Pro Gly  Thr Ser Glu
    2315                 2320                 2325

Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser
    2330                 2335                 2340

Glu Gly  Ser Ala Pro Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    2345                 2350                 2355

Ala Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    2360                 2365                 2370

Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr
    2375                 2380                 2385

Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser
    2390                 2395                 2400

Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr
    2405                 2410                 2415

Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    2420                 2425                 2430

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro
    2435                 2440                 2445

Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr
    2450                 2455                 2460
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Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu
    2465                 2470                 2475


Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    2480                 2485                 2490


Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Glu
    2495                 2500                 2505


Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser Pro Ala  Gly Ser Pro
    2510                 2515                 2520


Thr Ser  Thr Glu Glu Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr
    2525                 2530                 2535


Glu Glu  Gly Ser Pro Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly
    2540                 2545                 2550


Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Thr
    2555                 2560                 2565


Glu Pro  Ser Glu Gly Ser Ala  Pro
    2570                 2575


<210> 1558
<211> 2576
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
      polypeptide

<400> 1558
Met Gln Ile Glu Leu Ser Thr Cys Phe Phe Leu Cys Leu Leu Arg Phe
1               5                  10                  15


Cys Phe Ser Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
            20                  25                  30


Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
        35                  40                  45


Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
    50                  55                  60


Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
65                  70                  75                  80


Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
                85                  90                  95


Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
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            100                 105                 110

Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
        115                 120                 125

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
    130                 135                 140

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
145                 150                 155                 160

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
                165                 170                 175

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu
            180                 185                 190

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
        195                 200                 205

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
    210                 215                 220

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
225                 230                 235                 240

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
                245                 250                 255

Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            260                 265                 270

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
        275                 280                 285

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
    290                 295                 300

Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
305                 310                 315                 320

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
                325                 330                 335

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
            340                 345                 350

Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
        355                 360                 365

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
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370 375 380

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
385 390 395 400

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
405 410 415

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
420 425 430

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
435 440 445

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
450 455 460

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
465 470 475 480

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
485 490 495

Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
500 505 510

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
515 520 525

Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
530 535 540

Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
545 550 555 560

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
565 570 575

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
580 585 590

Ser Ala Pro Gln Ser Pro Arg Ser Phe Gln Gly Pro Glu Gly Pro Ser
595 600 605

Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
610 615 620

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
625 630 635 640

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys

645 650 655

Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro
660 665 670

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
675 680 685

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
690 695 700

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
705 710 715 720

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
725 730 735

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
740 745 750

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
755 760 765

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
770 775 780

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
785 790 795 800

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
805 810 815

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
820 825 830

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
835 840 845

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
850 855 860

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
865 870 875 880

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
885 890 895

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
900 905 910

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met

915 920 925

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
930 935 940

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
945 950 955 960

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
965 970 975

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
980 985 990

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
995 1000 1005

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly
1010 1015 1020

Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala
1025 1030 1035

Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu
1040 1045 1050

Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr
1055 1060 1065

Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile
1070 1075 1080

Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg
1085 1090 1095

Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro
1100 1105 1110

Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly
1115 1120 1125

Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser
1130 1135 1140

Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro
1145 1150 1155

Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln
1160 1165 1170

Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp

1175 1180 1185

Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu
1190 1195 1200

Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala
1205 1210 1215

Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu
1220 1225 1230

Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
1235 1240 1245

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly
1250 1255 1260

Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu
1265 1270 1275

Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro
1280 1285 1290

Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg
1295 1300 1305

Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr
1310 1315 1320

Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu
1325 1330 1335

Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn
1340 1345 1350

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
1355 1360 1365

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro
1370 1375 1380

Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
1385 1390 1395

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
1400 1405 1410

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
1415 1420 1425

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr

1430 1435 1440

Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
1445 1450 1455

Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
1460 1465 1470

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
1475 1480 1485

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro
1490 1495 1500

Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
1505 1510 1515

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
1520 1525 1530

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
1535 1540 1545

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro
1550 1555 1560

Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
1565 1570 1575

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
1580 1585 1590

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
1595 1600 1605

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala
1610 1615 1620

Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
1625 1630 1635

Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
1640 1645 1650

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
1655 1660 1665

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala
1670 1675 1680

Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly

1685 1690 1695

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
1700 1705 1710

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
1715 1720 1725

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro
1730 1735 1740

Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
1745 1750 1755

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
1760 1765 1770

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
1775 1780 1785

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr
1790 1795 1800

Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
1805 1810 1815

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
1820 1825 1830

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
1835 1840 1845

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala
1850 1855 1860

Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
1865 1870 1875

Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
1880 1885 1890

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
1895 1900 1905

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro
1910 1915 1920

Ser Glu Gly Ser Ala Pro Gln Ser Pro Arg Ser Phe Gln Lys Lys
1925 1930 1935

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr

1940 1945 1950

Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser
1955 1960 1965

Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr
1970 1975 1980

Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu
1985 1990 1995

His Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp
2000 2005 2010

Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser
2015 2020 2025

Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg Gln Gly
2030 2035 2040

Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr Lys Thr
2045 2050 2055

Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp Glu
2060 2065 2070

Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu
2075 2080 2085

Lys Asp Val His Ser Gly Leu Ile Gly Pro Leu Leu Val Cys His
2090 2095 2100

Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr Val Gln
2105 2110 2115

Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys Ser Trp
2120 2125 2130

Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro Cys Asn
2135 2140 2145

Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg Phe His
2150 2155 2160

Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu Val Met
2165 2170 2175

Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met Gly Ser
2180 2185 2190

Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val Phe Thr

2195 2200 2205

Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn Leu Tyr
2210 2215 2220

Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys Ala Gly
2225 2230 2235

Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His Ala Gly
2240 2245 2250

Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln Thr Pro
2255 2260 2265

Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile Thr Ala
2270 2275 2280

Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg Leu His
2285 2290 2295

Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro Phe Ser
2300 2305 2310

Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His Gly Ile
2315 2320 2325

Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr Ile Ser
2330 2335 2340

Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp Gln Thr
2345 2350 2355

Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe Gly Asn
2360 2365 2370

Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro Pro Ile
2375 2380 2385

Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser Ile Arg
2390 2395 2400

Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn Ser Cys
2405 2410 2415

Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp Ala Gln
2420 2425 2430

Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp Ser
2435 2440 2445

Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp

2450 2455 2460

Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe
2465 2470 2475

Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys
2480 2485 2490

Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser
2495 2500 2505

Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys
2510 2515 2520

Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val
2525 2530 2535

Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His
2540 2545 2550

Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu
2555 2560 2565

Gly Cys Glu Ala Gln Asp Leu Tyr
2570 2575

<210> 1559
<211> 2871
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1559
Met Gln Ile Glu Leu Ser Thr Cys Phe Phe Leu Cys Leu Leu Arg Phe
1 5 10 15

Cys Phe Ser Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
20 25 30

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
35 40 45

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
50 55 60

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
65 70 75 80

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
85 90 95

```
Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
            100                 105                 110

Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
        115                 120                 125

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
    130                 135                 140

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
145                 150                 155                 160

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
                165                 170                 175

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu
            180                 185                 190

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
        195                 200                 205

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
    210                 215                 220

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
225                 230                 235                 240

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
                245                 250                 255

Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
            260                 265                 270

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
        275                 280                 285

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
    290                 295                 300

Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
305                 310                 315                 320

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
                325                 330                 335

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
            340                 345                 350

Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
        355                 360                 365
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Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
    370                 375                 380

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
385                 390                 395                 400

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
                405                 410                 415

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
            420                 425                 430

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
        435                 440                 445

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
    450                 455                 460

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
465                 470                 475                 480

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
                485                 490                 495

Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
            500                 505                 510

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
        515                 520                 525

Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
    530                 535                 540

Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
545                 550                 555                 560

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
                565                 570                 575

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
            580                 585                 590

Ser Ala Pro Ile Glu Pro Arg Ser Pro Ser Gly Ser Pro Gly Ala Thr
        595                 600                 605

Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr Met Gln
    610                 615                 620

Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro Arg Val
625                 630                 635                 640
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Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys Thr Leu
                645                 650                 655

Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro Arg Pro
            660                 665                 670

Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val Tyr Asp
        675                 680                 685

Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val Ser Leu
    690                 695                 700

His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala Glu Tyr
705                 710                 715                 720

Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val Phe Pro
                725                 730                 735

Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn Gly Pro
            740                 745                 750

Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser His Val
        755                 760                 765

Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu Leu Val
    770                 775                 780

Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu His Lys
785                 790                 795                 800

Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp His Ser
                805                 810                 815

Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser Ala Arg
            820                 825                 830

Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg Ser Leu
        835                 840                 845

Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His Val Ile
    850                 855                 860

Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu Gly His
865                 870                 875                 880

Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile Ser Pro
                885                 890                 895

Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly Gln Phe
            900                 905                 910
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Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met Glu Ala
        915                 920                 925

Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg Met Lys
    930                 935                 940

Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp Ser Glu
945                 950                 955                 960

Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe Ile Gln
                965                 970                 975

Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His Tyr Ile
            980                 985                 990

Ala Ala Glu Glu Glu Asp Trp Asp  Tyr Ala Pro Leu Val  Leu Ala Pro
        995                 1000                1005

Asp Asp  Arg Ser Tyr Lys Ser  Gln Tyr Leu Asn Asn  Gly Pro Gln
    1010                1015                1020

Arg Ile  Gly Arg Lys Tyr Lys  Lys Val Arg Phe Met  Ala Tyr Thr
    1025                1030                1035

Asp Glu  Thr Phe Lys Thr Arg  Glu Ala Ile Gln His  Glu Ser Gly
    1040                1045                1050

Ile Leu  Gly Pro Leu Leu Tyr  Gly Glu Val Gly Asp  Thr Leu Leu
    1055                1060                1065

Ile Ile  Phe Lys Asn Gln Ala  Ser Arg Pro Tyr Asn  Ile Tyr Pro
    1070                1075                1080

His Gly  Ile Thr Asp Val Arg  Pro Leu Tyr Ser Arg  Arg Leu Pro
    1085                1090                1095

Lys Gly  Val Lys His Leu Lys  Asp Phe Pro Ile Leu  Pro Gly Glu
    1100                1105                1110

Ile Phe  Lys Tyr Lys Trp Thr  Val Thr Val Glu Asp  Gly Pro Thr
    1115                1120                1125

Lys Ser  Asp Pro Arg Cys Leu  Thr Arg Tyr Tyr Ser  Ser Phe Val
    1130                1135                1140

Asn Met  Glu Arg Asp Leu Ala  Ser Gly Leu Ile Gly  Pro Leu Leu
    1145                1150                1155

Ile Cys  Tyr Lys Glu Ser Val  Asp Gln Arg Gly Asn  Gln Ile Met
    1160                1165                1170
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Ser Asp  Lys Arg Asn Val Ile  Leu Phe Ser Val Phe  Asp Glu Asn
    1175                1180                1185

Arg Ser  Trp Tyr Leu Thr Glu  Asn Ile Gln Arg Phe  Leu Pro Asn
    1190                1195                1200

Pro Ala  Gly Val Gln Leu Glu  Asp Pro Glu Phe Gln  Ala Ser Asn
    1205                1210                1215

Ile Met  His Ser Ile Asn Gly  Tyr Val Phe Asp Ser  Leu Gln Leu
    1220                1225                1230

Ser Val  Cys Leu His Glu Val  Ala Tyr Trp Tyr Ile  Leu Ser Ile
    1235                1240                1245

Gly Ala  Gln Thr Asp Phe Leu  Ser Val Phe Phe Ser  Gly Tyr Thr
    1250                1255                1260

Phe Lys  His Lys Met Val Tyr  Glu Asp Thr Leu Thr  Leu Phe Pro
    1265                1270                1275

Phe Ser  Gly Glu Thr Val Phe  Met Ser Met Glu Asn  Pro Gly Leu
    1280                1285                1290

Trp Ile  Leu Gly Cys His Asn  Ser Asp Phe Arg Asn  Arg Gly Met
    1295                1300                1305

Thr Ala  Leu Leu Lys Val Ser  Ser Cys Asp Lys Asn  Thr Gly Asp
    1310                1315                1320

Tyr Tyr  Glu Asp Ser Tyr Glu  Asp Ile Ser Ala Tyr  Leu Leu Ser
    1325                1330                1335

Lys Asn  Asn Ala Ile Glu Pro  Arg Ser Phe Ser Gly  Thr Ser Glu
    1340                1345                1350

Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser
    1355                1360                1365

Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1370                1375                1380

Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly
    1385                1390                1395

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Thr
    1400                1405                1410

Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser Pro Ala  Gly Ser Pro
    1415                1420                1425
```

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Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1430                1435                1440

Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly
    1445                1450                1455

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Ser Pro Ala
    1460                1465                1470

Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro
    1475                1480                1485

Thr Ser  Thr Glu Glu Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    1490                1495                1500

Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    1505                1510                1515

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Glu
    1520                1525                1530

Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser
    1535                1540                1545

Gly Ser  Glu Thr Pro Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu
    1550                1555                1560

Thr Pro  Gly Ser Pro Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly
    1565                1570                1575

Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr
    1580                1585                1590

Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser Glu Pro  Ala Thr Ser
    1595                1600                1605

Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1610                1615                1620

Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gln
    1625                1630                1635

Ser Pro  Arg Ser Phe Gln Lys  Lys Thr Arg His Tyr  Phe Ile Ala
    1640                1645                1650

Ala Val  Glu Arg Leu Trp Asp  Tyr Gly Met Ser Ser  Ser Pro His
    1655                1660                1665

Val Leu  Arg Asn Arg Ala Gln  Ser Gly Ser Val Pro  Gln Phe Lys
    1670                1675                1680
```

```
Lys Val  Val Phe Gln Glu Phe  Thr Asp Gly Ser Phe  Thr Gln Pro
    1685                1690                1695

Leu Tyr  Arg Gly Glu Leu Asn  Glu His Leu Gly Leu  Leu Gly Pro
    1700                1705                1710

Tyr Ile  Arg Ala Glu Val Glu  Asp Asn Ile Met Val  Thr Phe Arg
    1715                1720                1725

Asn Gln  Ala Ser Arg Pro Tyr  Ser Phe Tyr Ser Ser  Leu Ile Ser
    1730                1735                1740

Tyr Glu  Glu Asp Gln Arg Gln  Gly Ala Glu Pro Arg  Lys Asn Phe
    1745                1750                1755

Val Lys  Pro Asn Glu Thr Lys  Thr Tyr Phe Trp Lys  Val Gln His
    1760                1765                1770

His Met  Ala Pro Thr Lys Asp  Glu Phe Asp Cys Lys  Ala Trp Ala
    1775                1780                1785

Tyr Phe  Ser Asp Val Asp Leu  Glu Lys Asp Val His  Ser Gly Leu
    1790                1795                1800

Ile Gly  Pro Leu Leu Val Cys  His Thr Asn Thr Leu  Asn Pro Ala
    1805                1810                1815

His Gly  Arg Gln Val Thr Val  Gln Glu Phe Ala Leu  Phe Phe Thr
    1820                1825                1830

Ile Phe  Asp Glu Thr Lys Ser  Trp Tyr Phe Thr Glu  Asn Met Glu
    1835                1840                1845

Arg Asn  Cys Arg Ala Pro Cys  Asn Ile Gln Met Glu  Asp Pro Thr
    1850                1855                1860

Phe Lys  Glu Asn Tyr Arg Phe  His Ala Ile Asn Gly  Tyr Ile Met
    1865                1870                1875

Asp Thr  Leu Pro Gly Leu Val  Met Ala Gln Asp Gln  Arg Ile Arg
    1880                1885                1890

Trp Tyr  Leu Leu Ser Met Gly  Ser Asn Glu Asn Ile  His Ser Ile
    1895                1900                1905

His Phe  Ser Gly His Val Phe  Thr Val Arg Lys Lys  Glu Glu Tyr
    1910                1915                1920

Lys Met  Ala Leu Tyr Asn Leu  Tyr Pro Gly Val Phe  Glu Thr Val
    1925                1930                1935
```

```
Glu Met  Leu Pro Ser Lys Ala  Gly Ile Trp Arg Val  Glu Cys Leu
    1940                1945                1950

Ile Gly  Glu His Leu His Ala  Gly Met Ser Thr Leu  Phe Leu Val
    1955                1960                1965

Tyr Ser  Asn Lys Cys Gln Thr  Pro Leu Gly Met Ala  Ser Gly His
    1970                1975                1980

Ile Arg  Asp Phe Gln Ile Thr  Ala Ser Gly Gln Tyr  Gly Gln Trp
    1985                1990                1995

Ala Pro  Lys Leu Ala Arg Leu  His Tyr Ser Gly Ser  Ile Asn Ala
    2000                2005                2010

Trp Ser  Thr Lys Glu Pro Phe  Ser Trp Ile Lys Val  Asp Leu Leu
    2015                2020                2025

Ala Pro  Met Ile Ile His Gly  Ile Lys Thr Gln Gly  Ala Arg Gln
    2030                2035                2040

Lys Phe  Ser Ser Leu Tyr Ile  Ser Gln Phe Ile Ile  Met Tyr Ser
    2045                2050                2055

Leu Asp  Gly Lys Lys Trp Gln  Thr Tyr Arg Gly Asn  Ser Thr Gly
    2060                2065                2070

Thr Leu  Met Val Phe Phe Gly  Asn Val Asp Ser Ser  Gly Ile Lys
    2075                2080                2085

His Asn  Ile Phe Asn Pro Pro  Ile Ile Ala Arg Tyr  Ile Arg Leu
    2090                2095                2100

His Pro  Thr His Tyr Ser Ile  Arg Ser Thr Leu Arg  Met Glu Leu
    2105                2110                2115

Met Gly  Cys Asp Leu Asn Ser  Cys Ser Met Pro Leu  Gly Met Glu
    2120                2125                2130

Ser Lys  Ala Ile Ser Asp Ala  Gln Ile Thr Ala Ser  Ser Tyr Phe
    2135                2140                2145

Thr Asn  Met Phe Ala Thr Trp  Ser Pro Ser Lys Ala  Arg Leu His
    2150                2155                2160

Leu Gln  Gly Arg Ser Asn Ala  Trp Arg Pro Gln Val  Asn Asn Pro
    2165                2170                2175

Lys Glu  Trp Leu Gln Val Asp  Phe Gln Lys Thr Met  Lys Val Thr
    2180                2185                2190
```

```
Gly Val  Thr Thr Gln Gly Val  Lys Ser Leu Leu Thr  Ser Met Tyr
    2195                 2200                 2205

Val Lys  Glu Phe Leu Ile Ser  Ser Ser Gln Asp Gly  His Gln Trp
    2210                 2215                 2220

Thr Leu  Phe Phe Gln Asn Gly  Lys Val Lys Val Phe  Gln Gly Asn
    2225                 2230                 2235

Gln Asp  Ser Phe Thr Pro Val  Val Asn Ser Leu Asp  Pro Pro Leu
    2240                 2245                 2250

Leu Thr  Arg Tyr Leu Arg Ile  His Pro Gln Ser Trp  Val His Gln
    2255                 2260                 2265

Ile Ala  Leu Arg Met Glu Val  Leu Gly Cys Glu Ala  Gln Asp Leu
    2270                 2275                 2280

Tyr Gly  Ser Pro Gly Ile Glu  Pro Arg Ser Pro Ser  Gly Ser Pro
    2285                 2290                 2295

Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Thr Ser  Glu Ser Ala
    2300                 2305                 2310

Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    2315                 2320                 2325

Ser Ala  Pro Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    2330                 2335                 2340

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    2345                 2350                 2355

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala
    2360                 2365                 2370

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    2375                 2380                 2385

Glu Thr  Pro Gly Ser Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro
    2390                 2395                 2400

Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser
    2405                 2410                 2415

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro
    2420                 2425                 2430

Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    2435                 2440                 2445
```

```
Ser Ala  Pro Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    2450                 2455                 2460

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    2465                 2470                 2475

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala
    2480                 2485                 2490

Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    2495                 2500                 2505

Ser Ala  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    2510                 2515                 2520

Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser
    2525                 2530                 2535

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Thr Glu Pro
    2540                 2545                 2550

Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    2555                 2560                 2565

Ser Gly  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    2570                 2575                 2580

Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser
    2585                 2590                 2595

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
    2600                 2605                 2610

Ser Gly  Ser Glu Thr Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    2615                 2620                 2625

Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    2630                 2635                 2640

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    2645                 2650                 2655

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Thr Glu Pro
    2660                 2665                 2670

Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    2675                 2680                 2685

Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    2690                 2695                 2700
```

```
Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser
    2705                 2710                 2715

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala
    2720                 2725                 2730

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    2735                 2740                 2745

Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    2750                 2755                 2760

Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser
    2765                 2770                 2775

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro
    2780                 2785                 2790

Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    2795                 2800                 2805

Ser Gly  Pro Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    2810                 2815                 2820

Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr Glu Glu  Gly Ser Pro
    2825                 2830                 2835

Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Thr Ser  Glu Ser Ala
    2840                 2845                 2850

Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    2855                 2860                 2865

Ser Ala  Pro
    2870


<210> 1560
<211> 2583
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
      polypeptide

<400> 1560
Met Gln Ile Glu Leu Ser Thr Cys Phe Phe Leu Cys Leu Leu Arg Phe
1               5                  10                  15

Cys Phe Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
            20                  25                  30

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
```

```
        35                  40                  45

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
    50                  55                  60

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
65                  70                  75                  80

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
                85                  90                  95

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
            100                 105                 110

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
        115                 120                 125

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
    130                 135                 140

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
145                 150                 155                 160

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
                165                 170                 175

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
            180                 185                 190

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
        195                 200                 205

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    210                 215                 220

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
225                 230                 235                 240

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
                245                 250                 255

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
            260                 265                 270

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
        275                 280                 285

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
    290                 295                 300

Thr Gly Ser Ile Glu Pro Arg Ser Pro Ser Gly Ser Pro Gly Ala Thr
```

```
305                 310                 315                 320

Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr Met Gln
                325                 330                 335

Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro Arg Val
            340                 345                 350

Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys Thr Leu
        355                 360                 365

Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro Arg Pro
    370                 375                 380

Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val Tyr Asp
385                 390                 395                 400

Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val Ser Leu
                405                 410                 415

His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala Glu Tyr
            420                 425                 430

Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val Phe Pro
        435                 440                 445

Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn Gly Pro
    450                 455                 460

Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser His Val
465                 470                 475                 480

Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu Leu Val
                485                 490                 495

Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu His Lys
            500                 505                 510

Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp His Ser
        515                 520                 525

Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser Ala Arg
    530                 535                 540

Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg Ser Leu
545                 550                 555                 560

Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His Val Ile
                565                 570                 575

Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu Gly His
```

580 585 590

Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile Ser Pro
595 600 605

Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly Gln Phe
610 615 620

Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met Glu Ala
625 630 635 640

Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg Met Lys
645 650 655

Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp Ser Glu
660 665 670

Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe Ile Gln
675 680 685

Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His Tyr Ile
690 695 700

Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu Ala Pro
705 710 715 720

Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg
725 730 735

Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu
740 745 750

Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly
755 760 765

Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys
770 775 780

Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp
785 790 795 800

Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu
805 810 815

Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr
820 825 830

Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr
835 840 845

Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly

```
    850                 855                 860

Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg
865                 870                 875                 880

Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val
                885                 890                 895

Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe
            900                 905                 910

Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala
        915                 920                 925

Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln
    930                 935                 940

Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile
945                 950                 955                 960

Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe
                965                 970                 975

Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser
            980                 985                 990

Gly Glu Thr Val Phe Met Ser Met  Glu Asn Pro Gly Leu  Trp Ile Leu
        995                 1000                1005

Gly Cys  His Asn Ser Asp Phe  Arg Asn Arg Gly Met  Thr Ala Leu
    1010                1015                1020

Leu Lys  Val Ser Ser Cys Asp  Lys Asn Thr Gly Asp  Tyr Tyr Glu
    1025                1030                1035

Asp Ser  Tyr Glu Asp Ile Ser  Ala Tyr Leu Leu Ser  Lys Asn Asn
    1040                1045                1050

Ala Ile  Glu Pro Arg Ser Phe  Ser Pro Gly Thr Pro  Gly Ser Gly
    1055                1060                1065

Thr Ala  Ser Ser Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    1070                1075                1080

Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro
    1085                1090                1095

Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr Gly Pro  Gly Ser Ser
    1100                1105                1110

Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser
```

1115 1120 1125

Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
1130 1135 1140

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
1145 1150 1155

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
1160 1165 1170

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr
1175 1180 1185

Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
1190 1195 1200

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
1205 1210 1215

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro
1220 1225 1230

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr
1235 1240 1245

Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
1250 1255 1260

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
1265 1270 1275

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
1280 1285 1290

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
1295 1300 1305

Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
1310 1315 1320

Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
1325 1330 1335

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
1340 1345 1350

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala
1355 1360 1365

Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr

1370 1375 1380

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
1385 1390 1395

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser
1400 1405 1410

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
1415 1420 1425

Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
1430 1435 1440

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
1445 1450 1455

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser
1460 1465 1470

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly
1475 1480 1485

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
1490 1495 1500

Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
1505 1510 1515

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
1520 1525 1530

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala
1535 1540 1545

Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
1550 1555 1560

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
1565 1570 1575

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser
1580 1585 1590

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala
1595 1600 1605

Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
1610 1615 1620

Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Gln

```
    1625                1630                1635

Ser Pro  Arg Ser Phe Gln Lys  Lys Thr Arg His Tyr  Phe Ile Ala
    1640                1645                1650

Ala Val  Glu Arg Leu Trp Asp  Tyr Gly Met Ser Ser  Ser Pro His
    1655                1660                1665

Val Leu  Arg Asn Arg Ala Gln  Ser Gly Ser Val Pro  Gln Phe Lys
    1670                1675                1680

Lys Val  Val Phe Gln Glu Phe  Thr Asp Gly Ser Phe  Thr Gln Pro
    1685                1690                1695

Leu Tyr  Arg Gly Glu Leu Asn  Glu His Leu Gly Leu  Leu Gly Pro
    1700                1705                1710

Tyr Ile  Arg Ala Glu Val Glu  Asp Asn Ile Met Val  Thr Phe Arg
    1715                1720                1725

Asn Gln  Ala Ser Arg Pro Tyr  Ser Phe Tyr Ser Ser  Leu Ile Ser
    1730                1735                1740

Tyr Glu  Glu Asp Gln Arg Gln  Gly Ala Glu Pro Arg  Lys Asn Phe
    1745                1750                1755

Val Lys  Pro Asn Glu Thr Lys  Thr Tyr Phe Trp Lys  Val Gln His
    1760                1765                1770

His Met  Ala Pro Thr Lys Asp  Glu Phe Asp Cys Lys  Ala Trp Ala
    1775                1780                1785

Tyr Phe  Ser Asp Val Asp Leu  Glu Lys Asp Val His  Ser Gly Leu
    1790                1795                1800

Ile Gly  Pro Leu Leu Val Cys  His Thr Asn Thr Leu  Asn Pro Ala
    1805                1810                1815

His Gly  Arg Gln Val Thr Val  Gln Glu Phe Ala Leu  Phe Phe Thr
    1820                1825                1830

Ile Phe  Asp Glu Thr Lys Ser  Trp Tyr Phe Thr Glu  Asn Met Glu
    1835                1840                1845

Arg Asn  Cys Arg Ala Pro Cys  Asn Ile Gln Met Glu  Asp Pro Thr
    1850                1855                1860

Phe Lys  Glu Asn Tyr Arg Phe  His Ala Ile Asn Gly  Tyr Ile Met
    1865                1870                1875

Asp Thr  Leu Pro Gly Leu Val  Met Ala Gln Asp Gln  Arg Ile Arg
```

1880 1885 1890

Trp Tyr Leu Leu Ser Met Gly Ser Asn Glu Asn Ile His Ser Ile
1895 1900 1905

His Phe Ser Gly His Val Phe Thr Val Arg Lys Lys Glu Glu Tyr
1910 1915 1920

Lys Met Ala Leu Tyr Asn Leu Tyr Pro Gly Val Phe Glu Thr Val
1925 1930 1935

Glu Met Leu Pro Ser Lys Ala Gly Ile Trp Arg Val Glu Cys Leu
1940 1945 1950

Ile Gly Glu His Leu His Ala Gly Met Ser Thr Leu Phe Leu Val
1955 1960 1965

Tyr Ser Asn Lys Cys Gln Thr Pro Leu Gly Met Ala Ser Gly His
1970 1975 1980

Ile Arg Asp Phe Gln Ile Thr Ala Ser Gly Gln Tyr Gly Gln Trp
1985 1990 1995

Ala Pro Lys Leu Ala Arg Leu His Tyr Ser Gly Ser Ile Asn Ala
2000 2005 2010

Trp Ser Thr Lys Glu Pro Phe Ser Trp Ile Lys Val Asp Leu Leu
2015 2020 2025

Ala Pro Met Ile Ile His Gly Ile Lys Thr Gln Gly Ala Arg Gln
2030 2035 2040

Lys Phe Ser Ser Leu Tyr Ile Ser Gln Phe Ile Ile Met Tyr Ser
2045 2050 2055

Leu Asp Gly Lys Lys Trp Gln Thr Tyr Arg Gly Asn Ser Thr Gly
2060 2065 2070

Thr Leu Met Val Phe Phe Gly Asn Val Asp Ser Ser Gly Ile Lys
2075 2080 2085

His Asn Ile Phe Asn Pro Pro Ile Ile Ala Arg Tyr Ile Arg Leu
2090 2095 2100

His Pro Thr His Tyr Ser Ile Arg Ser Thr Leu Arg Met Glu Leu
2105 2110 2115

Met Gly Cys Asp Leu Asn Ser Cys Ser Met Pro Leu Gly Met Glu
2120 2125 2130

Ser Lys Ala Ile Ser Asp Ala Gln Ile Thr Ala Ser Ser Tyr Phe

2135 2140 2145

Thr Asn Met Phe Ala Thr Trp Ser Pro Ser Lys Ala Arg Leu His
2150 2155 2160

Leu Gln Gly Arg Ser Asn Ala Trp Arg Pro Gln Val Asn Asn Pro
2165 2170 2175

Lys Glu Trp Leu Gln Val Asp Phe Gln Lys Thr Met Lys Val Thr
2180 2185 2190

Gly Val Thr Thr Gln Gly Val Lys Ser Leu Leu Thr Ser Met Tyr
2195 2200 2205

Val Lys Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly His Gln Trp
2210 2215 2220

Thr Leu Phe Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn
2225 2230 2235

Gln Asp Ser Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu
2240 2245 2250

Leu Thr Arg Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln
2255 2260 2265

Ile Ala Leu Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu
2270 2275 2280

Tyr Gly Ser Pro Gly Gln Ser Pro Arg Ser Phe Gln Pro Gly Ala
2285 2290 2295

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
2300 2305 2310

Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
2315 2320 2325

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
2330 2335 2340

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser
2345 2350 2355

Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser
2360 2365 2370

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
2375 2380 2385

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser

2390 2395 2400

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser
2405 2410 2415

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly
2420 2425 2430

Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
2435 2440 2445

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
2450 2455 2460

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser
2465 2470 2475

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly
2480 2485 2490

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
2495 2500 2505

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
2510 2515 2520

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala
2525 2530 2535

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser
2540 2545 2550

Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala
2555 2560 2565

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
2570 2575 2580

<210> 1561
<211> 2873
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1561
Met Gln Ile Glu Leu Ser Thr Cys Phe Phe Leu Cys Leu Leu Arg Phe
1 5 10 15

Cys Phe Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
20 25 30

```
Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
        35                  40                  45

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
    50                  55                  60

Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
65                  70                  75                  80

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro
                85                  90                  95

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
            100                 105                 110

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
        115                 120                 125

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
    130                 135                 140

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
145                 150                 155                 160

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                165                 170                 175

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
            180                 185                 190

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
        195                 200                 205

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    210                 215                 220

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
225                 230                 235                 240

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
                245                 250                 255

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
            260                 265                 270

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
        275                 280                 285

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
    290                 295                 300
```

```
Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
305                 310                 315                 320

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
                325                 330                 335

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
            340                 345                 350

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
        355                 360                 365

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
    370                 375                 380

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
385                 390                 395                 400

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                405                 410                 415

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
            420                 425                 430

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
        435                 440                 445

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
    450                 455                 460

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
465                 470                 475                 480

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
                485                 490                 495

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            500                 505                 510

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
        515                 520                 525

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
    530                 535                 540

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
545                 550                 555                 560

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
                565                 570                 575
```

```
Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
            580                 585                 590

Thr Gly Ser Gln Ser Pro Arg Ser Phe Gln Gly Ser Pro Gly Ala Thr
        595                 600                 605

Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr Met Gln
    610                 615                 620

Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro Arg Val
625                 630                 635                 640

Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys Thr Leu
                645                 650                 655

Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro Arg Pro
            660                 665                 670

Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val Tyr Asp
        675                 680                 685

Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val Ser Leu
    690                 695                 700

His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala Glu Tyr
705                 710                 715                 720

Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val Phe Pro
                725                 730                 735

Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn Gly Pro
            740                 745                 750

Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser His Val
        755                 760                 765

Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu Leu Val
    770                 775                 780

Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu His Lys
785                 790                 795                 800

Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp His Ser
                805                 810                 815

Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser Ala Arg
            820                 825                 830

Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg Ser Leu
        835                 840                 845
```

```
Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His Val Ile
    850                 855                 860

Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu Gly His
865                 870                 875                 880

Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile Ser Pro
                885                 890                 895

Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly Gln Phe
            900                 905                 910

Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met Glu Ala
        915                 920                 925

Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg Met Lys
    930                 935                 940

Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp Ser Glu
945                 950                 955                 960

Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe Ile Gln
                965                 970                 975

Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His Tyr Ile
            980                 985                 990

Ala Ala Glu Glu Glu Asp Trp Asp  Tyr Ala Pro Leu Val  Leu Ala Pro
        995                 1000                1005

Asp Asp  Arg Ser Tyr Lys Ser  Gln Tyr Leu Asn Asn  Gly Pro Gln
    1010                1015                1020

Arg Ile  Gly Arg Lys Tyr Lys  Lys Val Arg Phe Met  Ala Tyr Thr
    1025                1030                1035

Asp Glu  Thr Phe Lys Thr Arg  Glu Ala Ile Gln His  Glu Ser Gly
    1040                1045                1050

Ile Leu  Gly Pro Leu Leu Tyr  Gly Glu Val Gly Asp  Thr Leu Leu
    1055                1060                1065

Ile Ile  Phe Lys Asn Gln Ala  Ser Arg Pro Tyr Asn  Ile Tyr Pro
    1070                1075                1080

His Gly  Ile Thr Asp Val Arg  Pro Leu Tyr Ser Arg  Arg Leu Pro
    1085                1090                1095

Lys Gly  Val Lys His Leu Lys  Asp Phe Pro Ile Leu  Pro Gly Glu
    1100                1105                1110
```

```
Ile Phe  Lys Tyr Lys Trp Thr  Val Thr Val Glu Asp  Gly Pro Thr
    1115                1120                1125

Lys Ser  Asp Pro Arg Cys Leu  Thr Arg Tyr Tyr Ser  Ser Phe Val
    1130                1135                1140

Asn Met  Glu Arg Asp Leu Ala  Ser Gly Leu Ile Gly  Pro Leu Leu
    1145                1150                1155

Ile Cys  Tyr Lys Glu Ser Val  Asp Gln Arg Gly Asn  Gln Ile Met
    1160                1165                1170

Ser Asp  Lys Arg Asn Val Ile  Leu Phe Ser Val Phe  Asp Glu Asn
    1175                1180                1185

Arg Ser  Trp Tyr Leu Thr Glu  Asn Ile Gln Arg Phe  Leu Pro Asn
    1190                1195                1200

Pro Ala  Gly Val Gln Leu Glu  Asp Pro Glu Phe Gln  Ala Ser Asn
    1205                1210                1215

Ile Met  His Ser Ile Asn Gly  Tyr Val Phe Asp Ser  Leu Gln Leu
    1220                1225                1230

Ser Val  Cys Leu His Glu Val  Ala Tyr Trp Tyr Ile  Leu Ser Ile
    1235                1240                1245

Gly Ala  Gln Thr Asp Phe Leu  Ser Val Phe Phe Ser  Gly Tyr Thr
    1250                1255                1260

Phe Lys  His Lys Met Val Tyr  Glu Asp Thr Leu Thr  Leu Phe Pro
    1265                1270                1275

Phe Ser  Gly Glu Thr Val Phe  Met Ser Met Glu Asn  Pro Gly Leu
    1280                1285                1290

Trp Ile  Leu Gly Cys His Asn  Ser Asp Phe Arg Asn  Arg Gly Met
    1295                1300                1305

Thr Ala  Leu Leu Lys Val Ser  Ser Cys Asp Lys Asn  Thr Gly Asp
    1310                1315                1320

Tyr Tyr  Glu Asp Ser Tyr Glu  Asp Ile Ser Ala Tyr  Leu Leu Ser
    1325                1330                1335

Lys Asn  Asn Ala Ile Glu Pro  Arg Ser Phe Ser Gln  Asn Pro Gly
    1340                1345                1350

Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser Pro Gly  Ala Ser Pro
    1355                1360                1365
```

```
Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Thr Pro Gly  Ser Gly Thr
    1370                1375                1380

Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly
    1385                1390                1395

Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly
    1400                1405                1410

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Thr Pro Gly
    1415                1420                1425

Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser Ser Thr  Pro Ser Gly
    1430                1435                1440

Ala Thr  Gly Ser Pro Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly
    1445                1450                1455

Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr Gly Thr  Gly Pro Gly
    1460                1465                1470

Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly Pro Gly  Ala Ser Pro
    1475                1480                1485

Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Thr Pro Gly  Ser Gly Thr
    1490                1495                1500

Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly
    1505                1510                1515

Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr Gly Thr  Gly Pro Gly
    1520                1525                1530

Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly Pro Gly  Ala Ser Pro
    1535                1540                1545

Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1550                1555                1560

Ser Thr  Gly Ser Pro Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly
    1565                1570                1575

Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr Gly Thr  Gly Pro Gly
    1580                1585                1590

Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser Pro Gly  Ser Ser Pro
    1595                1600                1605

Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Thr Pro Gly  Ser Gly Thr
    1610                1615                1620
```

```
Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly
    1625                1630                1635

Ser Gln  Ser Pro Arg Ser Phe  Gln Lys Lys Thr Arg  His Tyr Phe
    1640                1645                1650

Ile Ala  Ala Val Glu Arg Leu  Trp Asp Tyr Gly Met  Ser Ser Ser
    1655                1660                1665

Pro His  Val Leu Arg Asn Arg  Ala Gln Ser Gly Ser  Val Pro Gln
    1670                1675                1680

Phe Lys  Lys Val Val Phe Gln  Glu Phe Thr Asp Gly  Ser Phe Thr
    1685                1690                1695

Gln Pro  Leu Tyr Arg Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu
    1700                1705                1710

Gly Pro  Tyr Ile Arg Ala Glu  Val Glu Asp Asn Ile  Met Val Thr
    1715                1720                1725

Phe Arg  Asn Gln Ala Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu
    1730                1735                1740

Ile Ser  Tyr Glu Glu Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys
    1745                1750                1755

Asn Phe  Val Lys Pro Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val
    1760                1765                1770

Gln His  His Met Ala Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala
    1775                1780                1785

Trp Ala  Tyr Phe Ser Asp Val  Asp Leu Glu Lys Asp  Val His Ser
    1790                1795                1800

Gly Leu  Ile Gly Pro Leu Leu  Val Cys His Thr Asn  Thr Leu Asn
    1805                1810                1815

Pro Ala  His Gly Arg Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe
    1820                1825                1830

Phe Thr  Ile Phe Asp Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn
    1835                1840                1845

Met Glu  Arg Asn Cys Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp
    1850                1855                1860

Pro Thr  Phe Lys Glu Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr
    1865                1870                1875
```

```
Ile Met  Asp Thr Leu Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg
    1880                 1885                 1890

Ile Arg  Trp Tyr Leu Leu Ser  Met Gly Ser Asn Glu  Asn Ile His
    1895                 1900                 1905

Ser Ile  His Phe Ser Gly His  Val Phe Thr Val Arg  Lys Lys Glu
    1910                 1915                 1920

Glu Tyr  Lys Met Ala Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu
    1925                 1930                 1935

Thr Val  Glu Met Leu Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu
    1940                 1945                 1950

Cys Leu  Ile Gly Glu His Leu  His Ala Gly Met Ser  Thr Leu Phe
    1955                 1960                 1965

Leu Val  Tyr Ser Asn Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser
    1970                 1975                 1980

Gly His  Ile Arg Asp Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly
    1985                 1990                 1995

Gln Trp  Ala Pro Lys Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile
    2000                 2005                 2010

Asn Ala  Trp Ser Thr Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp
    2015                 2020                 2025

Leu Leu  Ala Pro Met Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala
    2030                 2035                 2040

Arg Gln  Lys Phe Ser Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met
    2045                 2050                 2055

Tyr Ser  Leu Asp Gly Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser
    2060                 2065                 2070

Thr Gly  Thr Leu Met Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly
    2075                 2080                 2085

Ile Lys  His Asn Ile Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile
    2090                 2095                 2100

Arg Leu  His Pro Thr His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met
    2105                 2110                 2115

Glu Leu  Met Gly Cys Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly
    2120                 2125                 2130
```

```
Met Glu  Ser Lys Ala Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser
    2135                2140                2145

Tyr Phe  Thr Asn Met Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg
    2150                2155                2160

Leu His  Leu Gln Gly Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn
    2165                2170                2175

Asn Pro  Lys Glu Trp Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys
    2180                2185                2190

Val Thr  Gly Val Thr Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser
    2195                2200                2205

Met Tyr  Val Lys Glu Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His
    2210                2215                2220

Gln Trp  Thr Leu Phe Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln
    2225                2230                2235

Gly Asn  Gln Asp Ser Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro
    2240                2245                2250

Pro Leu  Leu Thr Arg Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val
    2255                2260                2265

His Gln  Ile Ala Leu Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln
    2270                2275                2280

Asp Leu  Tyr Gly Ser Pro Gly  Gln Ser Pro Arg Ser  Phe Gln Gly
    2285                2290                2295

Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Glu
    2300                2305                2310

Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser
    2315                2320                2325

Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr
    2330                2335                2340

Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    2345                2350                2355

Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Glu
    2360                2365                2370

Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser
    2375                2380                2385
```

```
Gly Ser  Glu Thr Pro Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu
    2390                 2395                 2400

Thr Pro  Gly Ser Pro Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly
    2405                 2410                 2415

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Thr
    2420                 2425                 2430

Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser
    2435                 2440                 2445

Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr
    2450                 2455                 2460

Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    2465                 2470                 2475

Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Glu
    2480                 2485                 2490

Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser
    2495                 2500                 2505

Glu Gly  Ser Ala Pro Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    2510                 2515                 2520

Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly
    2525                 2530                 2535

Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr
    2540                 2545                 2550

Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr Ser Glu  Ser Ala Thr
    2555                 2560                 2565

Pro Glu  Ser Gly Pro Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    2570                 2575                 2580

Gly Pro  Gly Ser Pro Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly
    2585                 2590                 2595

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro
    2600                 2605                 2610

Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr
    2615                 2620                 2625

Pro Glu  Ser Gly Pro Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    2630                 2635                 2640
```

```
Ala Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    2645                 2650                 2655

Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr
    2660                 2665                 2670

Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser
    2675                 2680                 2685

Glu Gly  Ser Ala Pro Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    2690                 2695                 2700

Ala Pro  Gly Ser Pro Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly
    2705                 2710                 2715

Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Glu
    2720                 2725                 2730

Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser
    2735                 2740                 2745

Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    2750                 2755                 2760

Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly
    2765                 2770                 2775

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Thr
    2780                 2785                 2790

Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr Ser Glu  Ser Ala Thr
    2795                 2800                 2805

Pro Glu  Ser Gly Pro Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr
    2810                 2815                 2820

Glu Glu  Gly Ser Pro Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly
    2825                 2830                 2835

Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Glu
    2840                 2845                 2850

Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser
    2855                 2860                 2865

Glu Gly  Ser Ala Pro
    2870
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<210> 1562
<211> 2572
<212> PRT

<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 1562
Met Gln Ile Glu Leu Ser Thr Cys Phe Phe Leu Cys Leu Leu Arg Phe
1 5 10 15

Cys Phe Ser Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser
20 25 30

Trp Asp Tyr Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg
35 40 45

Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val
50 55 60

Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile
65 70 75 80

Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln
85 90 95

Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser
100 105 110

His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser
115 120 125

Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp
130 135 140

Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu
145 150 155 160

Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser
165 170 175

Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile
180 185 190

Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr
195 200 205

Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly
210 215 220

Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp
225 230 235 240

Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr

```
                245                 250                 255

Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val
            260                 265                 270

Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile
        275                 280                 285

Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser
    290                 295                 300

Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met
305                 310                 315                 320

Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His
                325                 330                 335

Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro
            340                 345                 350

Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp
        355                 360                 365

Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser
    370                 375                 380

Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr
385                 390                 395                 400

Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro
                405                 410                 415

Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn
            420                 425                 430

Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met
        435                 440                 445

Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu
    450                 455                 460

Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu
465                 470                 475                 480

Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro
                485                 490                 495

His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys
            500                 505                 510

Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe
```

515 520 525

Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp
530 535 540

Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg
545 550 555 560

Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu
565 570 575

Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val
580 585 590

Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu
595 600 605

Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp
610 615 620

Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val
625 630 635 640

Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp
645 650 655

Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe
660 665 670

Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr
675 680 685

Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro
690 695 700

Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly
705 710 715 720

Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp
725 730 735

Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys
740 745 750

Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Pro Gly Thr Pro Gly Ser
755 760 765

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
770 775 780

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly

785 790 795 800

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr Pro
805 810 815

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
820 825 830

Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
835 840 845

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
850 855 860

Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
865 870 875 880

Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
885 890 895

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
900 905 910

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
915 920 925

Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
930 935 940

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly
945 950 955 960

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
965 970 975

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
980 985 990

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly
995 1000 1005

Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
1010 1015 1020

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
1025 1030 1035

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser
1040 1045 1050

Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser

```
    1055                1060                1065

Ala Ser  Thr Gly Thr Gly Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1070                1075                1080

Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    1085                1090                1095

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ala
    1100                1105                1110

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1115                1120                1125

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1130                1135                1140

Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    1145                1150                1155

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser
    1160                1165                1170

Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Thr  Pro Gly Ser
    1175                1180                1185

Gly Thr  Ala Ser Ser Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1190                1195                1200

Thr Gly  Ser Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1205                1210                1215

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser
    1220                1225                1230

Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Ser  Ser Pro Ser
    1235                1240                1245

Ala Ser  Thr Gly Thr Gly Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1250                1255                1260

Gly Thr  Gly Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    1265                1270                1275

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser
    1280                1285                1290

Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Ser  Ser Pro Ser
    1295                1300                1305

Ala Ser  Thr Gly Thr Gly Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
```

1310 1315 1320

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
1325 1330 1335

Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile
1340 1345 1350

Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro
1355 1360 1365

His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe
1370 1375 1380

Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln
1385 1390 1395

Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly
1400 1405 1410

Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe
1415 1420 1425

Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
1430 1435 1440

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn
1445 1450 1455

Phe Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln
1460 1465 1470

His His Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp
1475 1480 1485

Ala Tyr Phe Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly
1490 1495 1500

Leu Ile Gly Pro Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro
1505 1510 1515

Ala His Gly Arg Gln Val Thr Val Gln Glu Phe Ala Leu Phe Phe
1520 1525 1530

Thr Ile Phe Asp Glu Thr Lys Ser Trp Tyr Phe Thr Glu Asn Met
1535 1540 1545

Glu Arg Asn Cys Arg Ala Pro Cys Asn Ile Gln Met Glu Asp Pro
1550 1555 1560

Thr Phe Lys Glu Asn Tyr Arg Phe His Ala Ile Asn Gly Tyr Ile

1565 1570 1575

Met Asp Thr Leu Pro Gly Leu Val Met Ala Gln Asp Gln Arg Ile
1580 1585 1590

Arg Trp Tyr Leu Leu Ser Met Gly Ser Asn Glu Asn Ile His Ser
1595 1600 1605

Ile His Phe Ser Gly His Val Phe Thr Val Arg Lys Lys Glu Glu
1610 1615 1620

Tyr Lys Met Ala Leu Tyr Asn Leu Tyr Pro Gly Val Phe Glu Thr
1625 1630 1635

Val Glu Met Leu Pro Ser Lys Ala Gly Ile Trp Arg Val Glu Cys
1640 1645 1650

Leu Ile Gly Glu His Leu His Ala Gly Met Ser Thr Leu Phe Leu
1655 1660 1665

Val Tyr Ser Asn Lys Cys Gln Thr Pro Leu Gly Met Ala Ser Gly
1670 1675 1680

His Ile Arg Asp Phe Gln Ile Thr Ala Ser Gly Gln Tyr Gly Gln
1685 1690 1695

Trp Ala Pro Lys Leu Ala Arg Leu His Tyr Ser Gly Ser Ile Asn
1700 1705 1710

Ala Trp Ser Thr Lys Glu Pro Phe Ser Trp Ile Lys Val Asp Leu
1715 1720 1725

Leu Ala Pro Met Ile Ile His Gly Ile Lys Thr Gln Gly Ala Arg
1730 1735 1740

Gln Lys Phe Ser Ser Leu Tyr Ile Ser Gln Phe Ile Ile Met Tyr
1745 1750 1755

Ser Leu Asp Gly Lys Lys Trp Gln Thr Tyr Arg Gly Asn Ser Thr
1760 1765 1770

Gly Thr Leu Met Val Phe Phe Gly Asn Val Asp Ser Ser Gly Ile
1775 1780 1785

Lys His Asn Ile Phe Asn Pro Pro Ile Ile Ala Arg Tyr Ile Arg
1790 1795 1800

Leu His Pro Thr His Tyr Ser Ile Arg Ser Thr Leu Arg Met Glu
1805 1810 1815

Leu Met Gly Cys Asp Leu Asn Ser Cys Ser Met Pro Leu Gly Met

```
    1820                1825                1830

Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile Thr Ala  Ser Ser Tyr
    1835                1840                1845

Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro Ser Lys  Ala Arg Leu
    1850                1855                1860

His Leu  Gln Gly Arg Ser Asn  Ala Trp Arg Pro Gln  Val Asn Asn
    1865                1870                1875

Pro Lys  Glu Trp Leu Gln Val  Asp Phe Gln Lys Thr  Met Lys Val
    1880                1885                1890

Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser Leu Leu  Thr Ser Met
    1895                1900                1905

Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser Gln Asp  Gly His Gln
    1910                1915                1920

Trp Thr  Leu Phe Phe Gln Asn  Gly Lys Val Lys Val  Phe Gln Gly
    1925                1930                1935

Asn Gln  Asp Ser Phe Thr Pro  Val Val Asn Ser Leu  Asp Pro Pro
    1940                1945                1950

Leu Leu  Thr Arg Tyr Leu Arg  Ile His Pro Gln Ser  Trp Val His
    1955                1960                1965

Gln Ile  Ala Leu Arg Met Glu  Val Leu Gly Cys Glu  Ala Gln Asp
    1970                1975                1980

Leu Tyr  Gly Ser Pro Gly Gln  Ser Pro Arg Ser Phe  Gln Pro Gly
    1985                1990                1995

Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser Pro Gly  Ser Ser Thr
    2000                2005                2010

Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser
    2015                2020                2025

Thr Gly  Thr Gly Pro Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr
    2030                2035                2040

Gly Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly
    2045                2050                2055

Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Ala Ser Pro
    2060                2065                2070

Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
```

2075 2080 2085

Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
2090 2095 2100

Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
2105 2110 2115

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
2120 2125 2130

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
2135 2140 2145

Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr
2150 2155 2160

Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
2165 2170 2175

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
2180 2185 2190

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
2195 2200 2205

Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
2210 2215 2220

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
2225 2230 2235

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
2240 2245 2250

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
2255 2260 2265

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
2270 2275 2280

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
2285 2290 2295

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
2300 2305 2310

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
2315 2320 2325

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser

2330 2335 2340

Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
2345 2350 2355

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
2360 2365 2370

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
2375 2380 2385

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
2390 2395 2400

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
2405 2410 2415

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
2420 2425 2430

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
2435 2440 2445

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
2450 2455 2460

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
2465 2470 2475

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
2480 2485 2490

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser
2495 2500 2505

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
2510 2515 2520

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
2525 2530 2535

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
2540 2545 2550

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser
2555 2560 2565

Ser Thr Gly Ser
2570

<210> 1563

<211> 2297
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1563

```
Met Gln Ile Glu Leu Ser Thr Cys Phe Phe Leu Cys Leu Leu Arg Phe
1               5                  10                  15

Cys Phe Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
            20                  25                  30

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
        35                  40                  45

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
    50                  55                  60

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
65                  70                  75                  80

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
                85                  90                  95

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
            100                 105                 110

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
        115                 120                 125

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
    130                 135                 140

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
145                 150                 155                 160

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
                165                 170                 175

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
            180                 185                 190

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
        195                 200                 205

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    210                 215                 220

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
225                 230                 235                 240
```

```
Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
                245                 250                 255

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
            260                 265                 270

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
        275                 280                 285

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
    290                 295                 300

Thr Gly Ser Gln Ser Pro Arg Ser Phe Gln Gly Pro Ser Gly Pro Ala
305                 310                 315                 320

Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr Met
                325                 330                 335

Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro Arg
            340                 345                 350

Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys Thr
        355                 360                 365

Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro Arg
    370                 375                 380

Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val Tyr
385                 390                 395                 400

Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val Ser
                405                 410                 415

Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala Glu
            420                 425                 430

Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val Phe
        435                 440                 445

Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn Gly
    450                 455                 460

Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser His
465                 470                 475                 480

Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu Leu
                485                 490                 495

Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu His
            500                 505                 510
```

```
Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp His
        515                 520                 525

Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser Ala
    530                 535                 540

Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg Ser
545                 550                 555                 560

Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His Val
                565                 570                 575

Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu Gly
            580                 585                 590

His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile Ser
        595                 600                 605

Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly Gln
    610                 615                 620

Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met Glu
625                 630                 635                 640

Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg Met
                645                 650                 655

Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp Ser
            660                 665                 670

Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe Ile
        675                 680                 685

Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His Tyr
    690                 695                 700

Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu Ala
705                 710                 715                 720

Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln
                725                 730                 735

Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp
            740                 745                 750

Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu
        755                 760                 765

Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe
    770                 775                 780
```

```
Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr
785                 790                 795                 800

Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His
                805                 810                 815

Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp
            820                 825                 830

Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu
        835                 840                 845

Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser
    850                 855                 860

Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln
865                 870                 875                 880

Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser
                885                 890                 895

Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg
            900                 905                 910

Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln
        915                 920                 925

Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu
    930                 935                 940

Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser
945                 950                 955                 960

Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr
                965                 970                 975

Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe
            980                 985                 990

Ser Gly Glu Thr Val Phe Met Ser  Met Glu Asn Pro Gly  Leu Trp Ile
        995                 1000                1005

Leu Gly  Cys His Asn Ser Asp  Phe Arg Asn Arg Gly  Met Thr Ala
    1010                1015                1020

Leu Leu  Lys Val Ser Ser Cys  Asp Lys Asn Thr Gly  Asp Tyr Tyr
    1025                1030                1035

Glu Asp  Ser Tyr Glu Asp Ile  Ser Ala Tyr Leu Leu  Ser Lys Asn
    1040                1045                1050
```

```
Asn Ala  Ile Glu Pro Arg Ser  Phe Ser Pro Gly Ala  Ser Pro Gly
    1055                 1060                 1065

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1070                 1075                 1080

Thr Gly  Ser Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1085                 1090                 1095

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Thr
    1100                 1105                 1110

Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro
    1115                 1120                 1125

Ser Gly  Ala Thr Gly Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala
    1130                 1135                 1140

Ser Ser  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1145                 1150                 1155

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser
    1160                 1165                 1170

Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro Gly Ser  Ser Pro Ser
    1175                 1180                 1185

Ala Ser  Thr Gly Thr Gly Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1190                 1195                 1200

Thr Gly  Ser Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1205                 1210                 1215

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser
    1220                 1225                 1230

Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro Gly Ser  Ser Pro Ser
    1235                 1240                 1245

Ala Ser  Thr Gly Thr Gly Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1250                 1255                 1260

Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    1265                 1270                 1275

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser
    1280                 1285                 1290

Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro Gly Ala  Ser Pro Gly
    1295                 1300                 1305
```

```
Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1310                 1315                 1320

Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1325                 1330                 1335

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Gln Ser Pro
    1340                 1345                 1350

Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr Phe Ile  Ala Ala Val
    1355                 1360                 1365

Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser Ser Pro  His Val Leu
    1370                 1375                 1380

Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro Gln Phe  Lys Lys Val
    1385                 1390                 1395

Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe Thr Gln  Pro Leu Tyr
    1400                 1405                 1410

Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu Leu Gly  Pro Tyr Ile
    1415                 1420                 1425

Arg Ala  Glu Val Glu Asp Asn  Ile Met Val Thr Phe  Arg Asn Gln
    1430                 1435                 1440

Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser Leu Ile  Ser Tyr Glu
    1445                 1450                 1455

Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg Lys Asn  Phe Val Lys
    1460                 1465                 1470

Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys Val Gln  His His Met
    1475                 1480                 1485

Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys Ala Trp  Ala Tyr Phe
    1490                 1495                 1500

Ser Asp  Val Asp Leu Glu Lys  Asp Val His Ser Gly  Leu Ile Gly
    1505                 1510                 1515

Pro Leu  Leu Val Cys His Thr  Asn Thr Leu Asn Pro  Ala His Gly
    1520                 1525                 1530

Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu Phe Phe  Thr Ile Phe
    1535                 1540                 1545

Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu Asn Met  Glu Arg Asn
    1550                 1555                 1560
```

```
Cys Arg  Ala Pro Cys Asn Ile  Gln Met Glu Asp Pro  Thr Phe Lys
    1565                1570                1575

Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly Tyr Ile  Met Asp Thr
    1580                1585                1590

Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln Arg Ile  Arg Trp Tyr
    1595                1600                1605

Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile His Ser  Ile His Phe
    1610                1615                1620

Ser Gly  His Val Phe Thr Val  Arg Lys Lys Glu Glu  Tyr Lys Met
    1625                1630                1635

Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe Glu Thr  Val Glu Met
    1640                1645                1650

Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val Glu Cys  Leu Ile Gly
    1655                1660                1665

Glu His  Leu His Ala Gly Met  Ser Thr Leu Phe Leu  Val Tyr Ser
    1670                1675                1680

Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala Ser Gly  His Ile Arg
    1685                1690                1695

Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr Gly Gln  Trp Ala Pro
    1700                1705                1710

Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser Ile Asn  Ala Trp Ser
    1715                1720                1725

Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val Asp Leu  Leu Ala Pro
    1730                1735                1740

Met Ile  Ile His Gly Ile Lys  Thr Gln Gly Ala Arg  Gln Lys Phe
    1745                1750                1755

Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile Met Tyr  Ser Leu Asp
    1760                1765                1770

Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn Ser Thr  Gly Thr Leu
    1775                1780                1785

Met Val  Phe Phe Gly Asn Val  Asp Ser Ser Gly Ile  Lys His Asn
    1790                1795                1800

Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr Ile Arg  Leu His Pro
    1805                1810                1815
```

```
Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg Met Glu  Leu Met Gly
    1820                1825                1830

Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu Gly Met  Glu Ser Lys
    1835                1840                1845

Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser Ser Tyr  Phe Thr Asn
    1850                1855                1860

Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala Arg Leu  His Leu Gln
    1865                1870                1875

Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val Asn Asn  Pro Lys Glu
    1880                1885                1890

Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met Lys Val  Thr Gly Val
    1895                1900                1905

Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr Ser Met  Tyr Val Lys
    1910                1915                1920

Glu Phe  Leu Ile Ser Ser Ser  Gln Asp Gly His Gln  Trp Thr Leu
    1925                1930                1935

Phe Phe  Gln Asn Gly Lys Val  Lys Val Phe Gln Gly  Asn Gln Asp
    1940                1945                1950

Ser Phe  Thr Pro Val Val Asn  Ser Leu Asp Pro Pro  Leu Leu Thr
    1955                1960                1965

Arg Tyr  Leu Arg Ile His Pro  Gln Ser Trp Val His  Gln Ile Ala
    1970                1975                1980

Leu Arg  Met Glu Val Leu Gly  Cys Glu Ala Gln Asp  Leu Tyr Gly
    1985                1990                1995

Pro Ser  Gly Pro Gln Ser Pro  Arg Ser Phe Gln Gly  Thr Ser Glu
    2000                2005                2010

Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser
    2015                2020                2025

Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    2030                2035                2040

Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly
    2045                2050                2055

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Thr
    2060                2065                2070
```

```
Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser Pro Ala  Gly Ser Pro
    2075                 2080                 2085

Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    2090                 2095                 2100

Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly
    2105                 2110                 2115

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Ser Pro Ala
    2120                 2125                 2130

Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro
    2135                 2140                 2145

Thr Ser  Thr Glu Glu Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    2150                 2155                 2160

Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    2165                 2170                 2175

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Glu
    2180                 2185                 2190

Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser
    2195                 2200                 2205

Gly Ser  Glu Thr Pro Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu
    2210                 2215                 2220

Thr Pro  Gly Ser Pro Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly
    2225                 2230                 2235

Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr
    2240                 2245                 2250

Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser Glu Pro  Ala Thr Ser
    2255                 2260                 2265

Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    2270                 2275                 2280

Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro
    2285                 2290                 2295
```

<210> 1564
<211> 2572
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic

polypeptide

<400> 1564

```
Met Gln Ile Glu Leu Ser Thr Cys Phe Phe Leu Cys Leu Leu Arg Phe
1               5                  10                  15

Cys Phe Ser Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
            20                  25                  30

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
        35                  40                  45

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
    50                  55                  60

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
65                  70                  75                  80

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
                85                  90                  95

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
            100                 105                 110

Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
        115                 120                 125

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
    130                 135                 140

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
145                 150                 155                 160

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
                165                 170                 175

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu
            180                 185                 190

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
        195                 200                 205

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
    210                 215                 220

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
225                 230                 235                 240

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
                245                 250                 255

Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
```

```
            260                 265                 270

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
        275                 280                 285

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
    290                 295                 300

Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
305                 310                 315                 320

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
                325                 330                 335

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
            340                 345                 350

Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
        355                 360                 365

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
    370                 375                 380

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
385                 390                 395                 400

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
                405                 410                 415

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
            420                 425                 430

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
        435                 440                 445

Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
    450                 455                 460

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser
465                 470                 475                 480

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
                485                 490                 495

Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
            500                 505                 510

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser
        515                 520                 525

Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
```

530 535 540

Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
545 550 555 560

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
565 570 575

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
580 585 590

Ser Ala Pro Gln Ser Pro Arg Ser Phe Gln Gly Ser Pro Gly Ala Thr
595 600 605

Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr Met Gln
610 615 620

Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro Arg Val
625 630 635 640

Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys Thr Leu
645 650 655

Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro Arg Pro
660 665 670

Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val Tyr Asp
675 680 685

Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val Ser Leu
690 695 700

His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala Glu Tyr
705 710 715 720

Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val Phe Pro
725 730 735

Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn Gly Pro
740 745 750

Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser His Val
755 760 765

Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu Leu Val
770 775 780

Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu His Lys
785 790 795 800

Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp His Ser

805 810 815

Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser Ala Arg
820 825 830

Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg Ser Leu
835 840 845

Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His Val Ile
850 855 860

Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu Gly His
865 870 875 880

Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile Ser Pro
885 890 895

Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly Gln Phe
900 905 910

Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met Glu Ala
915 920 925

Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg Met Lys
930 935 940

Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp Ser Glu
945 950 955 960

Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe Ile Gln
965 970 975

Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His Tyr Ile
980 985 990

Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu Ala Pro
995 1000 1005

Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln
1010 1015 1020

Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
1025 1030 1035

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly
1040 1045 1050

Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu
1055 1060 1065

Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro

1070 1075 1080

His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro
1085 1090 1095

Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu
1100 1105 1110

Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr
1115 1120 1125

Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val
1130 1135 1140

Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu
1145 1150 1155

Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met
1160 1165 1170

Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn
1175 1180 1185

Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn
1190 1195 1200

Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn
1205 1210 1215

Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu
1220 1225 1230

Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile
1235 1240 1245

Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr
1250 1255 1260

Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
1265 1270 1275

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu
1280 1285 1290

Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met
1295 1300 1305

Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp
1310 1315 1320

Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser

```
    1325                1330                1335

Lys Asn  Asn Ala Ile Glu Pro  Arg Ser Phe Ser Pro  Gly Thr Pro
    1340                1345                1350

Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly Ser Ser  Thr Pro Ser
    1355                1360                1365

Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Pro Ser Ala  Ser Thr Gly
    1370                1375                1380

Thr Gly  Pro Gly Ser Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro
    1385                1390                1395

Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser
    1400                1405                1410

Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    1415                1420                1425

Ser Ser  Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
    1430                1435                1440

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1445                1450                1455

Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser Ser Pro  Gly Ala Ser
    1460                1465                1470

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    1475                1480                1485

Ser Ser  Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
    1490                1495                1500

Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro
    1505                1510                1515

Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser Ser Pro  Gly Ala Ser
    1520                1525                1530

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    1535                1540                1545

Ser Ser  Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
    1550                1555                1560

Gly Ser  Pro Gly Ser Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro
    1565                1570                1575

Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Gly Ala Ser
```

1580 1585 1590

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly
1595 1600 1605

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
1610 1615 1620

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
1625 1630 1635

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
1640 1645 1650

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr
1655 1660 1665

Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
1670 1675 1680

Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
1685 1690 1695

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
1700 1705 1710

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
1715 1720 1725

Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
1730 1735 1740

Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
1745 1750 1755

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro
1760 1765 1770

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser
1775 1780 1785

Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
1790 1795 1800

Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
1805 1810 1815

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
1820 1825 1830

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala

```
    1835                1840                1845

Ser Thr  Gly Thr Gly Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
    1850                1855                1860

Gly Ser  Pro Gly Thr Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro
    1865                1870                1875

Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser
    1880                1885                1890

Pro Ser  Ala Ser Thr Gly Thr  Gly Pro Gly Ser Ser  Pro Ser Ala
    1895                1900                1905

Ser Thr  Gly Thr Gly Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
    1910                1915                1920

Gly Ser  Gln Ser Pro Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr
    1925                1930                1935

Phe Ile  Ala Ala Val Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser
    1940                1945                1950

Ser Pro  His Val Leu Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro
    1955                1960                1965

Gln Phe  Lys Lys Val Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe
    1970                1975                1980

Thr Gln  Pro Leu Tyr Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu
    1985                1990                1995

Leu Gly  Pro Tyr Ile Arg Ala  Glu Val Glu Asp Asn  Ile Met Val
    2000                2005                2010

Thr Phe  Arg Asn Gln Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser
    2015                2020                2025

Leu Ile  Ser Tyr Glu Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg
    2030                2035                2040

Lys Asn  Phe Val Lys Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys
    2045                2050                2055

Val Gln  His His Met Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys
    2060                2065                2070

Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu Glu Lys  Asp Val His
    2075                2080                2085

Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys His Thr  Asn Thr Leu
```

2090 2095 2100

Asn Pro Ala His Gly Arg Gln Val Thr Val Gln Glu Phe Ala Leu
2105 2110 2115

Phe Phe Thr Ile Phe Asp Glu Thr Lys Ser Trp Tyr Phe Thr Glu
2120 2125 2130

Asn Met Glu Arg Asn Cys Arg Ala Pro Cys Asn Ile Gln Met Glu
2135 2140 2145

Asp Pro Thr Phe Lys Glu Asn Tyr Arg Phe His Ala Ile Asn Gly
2150 2155 2160

Tyr Ile Met Asp Thr Leu Pro Gly Leu Val Met Ala Gln Asp Gln
2165 2170 2175

Arg Ile Arg Trp Tyr Leu Leu Ser Met Gly Ser Asn Glu Asn Ile
2180 2185 2190

His Ser Ile His Phe Ser Gly His Val Phe Thr Val Arg Lys Lys
2195 2200 2205

Glu Glu Tyr Lys Met Ala Leu Tyr Asn Leu Tyr Pro Gly Val Phe
2210 2215 2220

Glu Thr Val Glu Met Leu Pro Ser Lys Ala Gly Ile Trp Arg Val
2225 2230 2235

Glu Cys Leu Ile Gly Glu His Leu His Ala Gly Met Ser Thr Leu
2240 2245 2250

Phe Leu Val Tyr Ser Asn Lys Cys Gln Thr Pro Leu Gly Met Ala
2255 2260 2265

Ser Gly His Ile Arg Asp Phe Gln Ile Thr Ala Ser Gly Gln Tyr
2270 2275 2280

Gly Gln Trp Ala Pro Lys Leu Ala Arg Leu His Tyr Ser Gly Ser
2285 2290 2295

Ile Asn Ala Trp Ser Thr Lys Glu Pro Phe Ser Trp Ile Lys Val
2300 2305 2310

Asp Leu Leu Ala Pro Met Ile Ile His Gly Ile Lys Thr Gln Gly
2315 2320 2325

Ala Arg Gln Lys Phe Ser Ser Leu Tyr Ile Ser Gln Phe Ile Ile
2330 2335 2340

Met Tyr Ser Leu Asp Gly Lys Lys Trp Gln Thr Tyr Arg Gly Asn

2345 2350 2355

Ser Thr Gly Thr Leu Met Val Phe Phe Gly Asn Val Asp Ser Ser
2360 2365 2370

Gly Ile Lys His Asn Ile Phe Asn Pro Pro Ile Ile Ala Arg Tyr
2375 2380 2385

Ile Arg Leu His Pro Thr His Tyr Ser Ile Arg Ser Thr Leu Arg
2390 2395 2400

Met Glu Leu Met Gly Cys Asp Leu Asn Ser Cys Ser Met Pro Leu
2405 2410 2415

Gly Met Glu Ser Lys Ala Ile Ser Asp Ala Gln Ile Thr Ala Ser
2420 2425 2430

Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp Ser Pro Ser Lys Ala
2435 2440 2445

Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg Pro Gln Val
2450 2455 2460

Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln Lys Thr Met
2465 2470 2475

Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser Leu Leu Thr
2480 2485 2490

Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly
2495 2500 2505

His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val Lys Val Phe
2510 2515 2520

Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn Ser Leu Asp
2525 2530 2535

Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro Gln Ser Trp
2540 2545 2550

Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly Cys Glu Ala
2555 2560 2565

Gln Asp Leu Tyr
2570

<210> 1565
<211> 2360
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1565

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Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1               5                  10                  15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Ser
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255
```

```
Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Lys
        355                 360                 365

Leu Thr Arg Ala Glu Thr Gly Glu Pro Ser Glu Gly Ser Ala Pro Gly
    370                 375                 380

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
385                 390                 395                 400

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
                405                 410                 415

Gly Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp
            420                 425                 430

Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu
        435                 440                 445

Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn
    450                 455                 460

Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala
465                 470                 475                 480

Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser
                485                 490                 495

Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu
            500                 505                 510

Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His
        515                 520                 525
```

```
Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly
    530                 535                 540

Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys
545                 550                 555                 560

Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro
                565                 570                 575

Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp
            580                 585                 590

Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser
        595                 600                 605

Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile
    610                 615                 620

Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn
625                 630                 635                 640

Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro
                645                 650                 655

Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe
            660                 665                 670

Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr
        675                 680                 685

Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser
    690                 695                 700

Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu
705                 710                 715                 720

Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly
                725                 730                 735

Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met
            740                 745                 750

Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr
        755                 760                 765

Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn
    770                 775                 780

Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys
785                 790                 795                 800
```

```
Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu
                805                 810                 815

Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp
            820                 825                 830

Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln
        835                 840                 845

Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp
    850                 855                 860

Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser
865                 870                 875                 880

Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp
                885                 890                 895

Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu
            900                 905                 910

Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met
        915                 920                 925

Val Thr Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser
    930                 935                 940

Leu Ile Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys
945                 950                 955                 960

Asn Phe Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln
                965                 970                 975

His His Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala
            980                 985                 990

Tyr Phe Ser Asp Val Asp Leu Glu  Lys Asp Val His Ser  Gly Leu Ile
        995                 1000                1005

Gly Pro  Leu Leu Val Cys His  Thr Asn Thr Leu Asn  Pro Ala His
    1010                1015                1020

Gly Arg  Gln Val Thr Val Gln  Glu Phe Ala Leu Phe  Phe Thr Ile
    1025                1030                1035

Phe Asp  Glu Thr Lys Ser Trp  Tyr Phe Thr Glu Asn  Met Glu Arg
    1040                1045                1050

Asn Cys  Arg Ala Pro Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe
    1055                1060                1065
```

```
Lys Glu  Asn Tyr Arg Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp
    1070                1075                1080

Thr Leu  Pro Gly Leu Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp
    1085                1090                1095

Tyr Leu  Leu Ser Met Gly Ser  Asn Glu Asn Ile His  Ser Ile His
    1100                1105                1110

Phe Ser  Gly His Val Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys
    1115                1120                1125

Met Ala  Leu Tyr Asn Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu
    1130                1135                1140

Met Leu  Pro Ser Lys Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile
    1145                1150                1155

Gly Glu  His Leu His Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr
    1160                1165                1170

Ser Asn  Lys Cys Gln Thr Pro  Leu Gly Met Ala Ser  Gly His Ile
    1175                1180                1185

Arg Asp  Phe Gln Ile Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala
    1190                1195                1200

Pro Lys  Leu Ala Arg Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp
    1205                1210                1215

Ser Thr  Lys Glu Pro Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala
    1220                1225                1230

Pro Met  Ile Ile His Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys
    1235                1240                1245

Phe Ser  Ser Leu Tyr Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu
    1250                1255                1260

Asp Gly  Lys Lys Trp Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr
    1265                1270                1275

Leu Met  Val Phe Phe Gly Asn  Val Asp Ser Ser Gly  Ile Lys His
    1280                1285                1290

Asn Ile  Phe Asn Pro Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His
    1295                1300                1305

Pro Thr  His Tyr Ser Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met
    1310                1315                1320
```

```
Gly Cys  Asp Leu Asn Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser
    1325                1330                1335

Lys Ala  Ile Ser Asp Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr
    1340                1345                1350

Asn Met  Phe Ala Thr Trp Thr  Pro Ser Lys Ala Arg  Leu His Leu
    1355                1360                1365

Gln Gly  Arg Ser Asn Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys
    1370                1375                1380

Glu Trp  Leu Gln Val Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly
    1385                1390                1395

Val Thr  Thr Gln Gly Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val
    1400                1405                1410

Lys Glu  Phe Leu Ile Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr
    1415                1420                1425

Leu Phe  Phe Gln Asn Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln
    1430                1435                1440

Asp Ser  Phe Thr Pro Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu
    1445                1450                1455

Thr Arg  Tyr Leu Arg Ile His  Pro Gln Ser Trp Val  His Gln Ile
    1460                1465                1470

Ala Leu  Arg Met Glu Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr
    1475                1480                1485

Lys Leu  Thr Arg Ala Glu Thr  Gly Gly Ser Pro Ala  Gly Ser Pro
    1490                1495                1500

Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1505                1510                1515

Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    1520                1525                1530

Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr
    1535                1540                1545

Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser
    1550                1555                1560

Glu Gly  Ser Ala Pro Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1565                1570                1575
```

```
Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly
    1580                 1585                 1590

Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr Pro Gly  Ser Pro Ala
    1595                 1600                 1605

Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr Ser Glu  Ser Ala Thr
    1610                 1615                 1620

Pro Glu  Ser Gly Pro Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    1625                 1630                 1635

Ala Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    1640                 1645                 1650

Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr
    1655                 1660                 1665

Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser
    1670                 1675                 1680

Glu Gly  Ser Ala Pro Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1685                 1690                 1695

Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    1700                 1705                 1710

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro
    1715                 1720                 1725

Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr Ser Thr  Glu Pro Ser
    1730                 1735                 1740

Glu Gly  Ser Ala Pro Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    1745                 1750                 1755

Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    1760                 1765                 1770

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Ser Pro Ala
    1775                 1780                 1785

Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr Ser Glu  Ser Ala Thr
    1790                 1795                 1800

Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu
    1805                 1810                 1815

Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    1820                 1825                 1830
```

```
Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr
    1835                 1840                 1845

Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser
    1850                 1855                 1860

Glu Gly  Ser Ala Pro Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    1865                 1870                 1875

Ala Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    1880                 1885                 1890

Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala Pro Gly  Ser Pro Ala
    1895                 1900                 1905

Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr Ser Thr  Glu Pro Ser
    1910                 1915                 1920

Glu Gly  Ser Ala Pro Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1925                 1930                 1935

Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly
    1940                 1945                 1950

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro
    1955                 1960                 1965

Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr
    1970                 1975                 1980

Pro Glu  Ser Gly Pro Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    1985                 1990                 1995

Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    2000                 2005                 2010

Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala
    2015                 2020                 2025

Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro
    2030                 2035                 2040

Thr Ser  Thr Glu Glu Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    2045                 2050                 2055

Gly Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    2060                 2065                 2070

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro
    2075                 2080                 2085
```

```
Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr
    2090                 2095                 2100

Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu
    2105                 2110                 2115

Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    2120                 2125                 2130

Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala Pro Gly  Ser Pro Ala
    2135                 2140                 2145

Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr Ser Glu  Ser Ala Thr
    2150                 2155                 2160

Pro Glu  Ser Gly Pro Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu
    2165                 2170                 2175

Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    2180                 2185                 2190

Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala
    2195                 2200                 2205

Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr Ser Thr  Glu Pro Ser
    2210                 2215                 2220

Glu Gly  Ser Ala Pro Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    2225                 2230                 2235

Gly Pro  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    2240                 2245                 2250

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro
    2255                 2260                 2265

Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Ser Glu Pro  Ala Thr Ser
    2270                 2275                 2280

Gly Ser  Glu Thr Pro Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr
    2285                 2290                 2295

Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    2300                 2305                 2310

Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala Pro Gly  Ser Glu Pro
    2315                 2320                 2325

Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr
    2330                 2335                 2340
```

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
2345 2350 2355

Ala Pro
2360

<210> 1566
<211> 2615
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1566
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp

195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile

```
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Leu Gly Arg Ile Val Gly
```

740 745 750

Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
755 760 765

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly
770 775 780

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
785 790 795 800

Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser
805 810 815

Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly
820 825 830

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
835 840 845

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser
850 855 860

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala
865 870 875 880

Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
885 890 895

Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser
900 905 910

Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala
915 920 925

Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr
930 935 940

Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
945 950 955 960

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser
965 970 975

Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr
980 985 990

Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser
995 1000 1005

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser

1010 1015 1020

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
1025 1030 1035

Thr Gly Ser Gly Pro Leu Gly Arg Ile Val Gly Gly Pro Pro Val
1040 1045 1050

Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser
1055 1060 1065

Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met
1070 1075 1080

Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser
1085 1090 1095

Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala
1100 1105 1110

Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val
1115 1120 1125

Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys
1130 1135 1140

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu
1145 1150 1155

Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr
1160 1165 1170

Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn
1175 1180 1185

Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr
1190 1195 1200

Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val
1205 1210 1215

Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
1220 1225 1230

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr
1235 1240 1245

Phe Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile
1250 1255 1260

Gly Pro Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His

1265 1270 1275

Gly Arg Gln Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile
1280 1285 1290

Phe Asp Glu Thr Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg
1295 1300 1305

Asn Cys Arg Ala Pro Cys Asn Ile Gln Met Glu Asp Pro Thr Phe
1310 1315 1320

Lys Glu Asn Tyr Arg Phe His Ala Ile Asn Gly Tyr Ile Met Asp
1325 1330 1335

Thr Leu Pro Gly Leu Val Met Ala Gln Asp Gln Arg Ile Arg Trp
1340 1345 1350

Tyr Leu Leu Ser Met Gly Ser Asn Glu Asn Ile His Ser Ile His
1355 1360 1365

Phe Ser Gly His Val Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys
1370 1375 1380

Met Ala Leu Tyr Asn Leu Tyr Pro Gly Val Phe Glu Thr Val Glu
1385 1390 1395

Met Leu Pro Ser Lys Ala Gly Ile Trp Arg Val Glu Cys Leu Ile
1400 1405 1410

Gly Glu His Leu His Ala Gly Met Ser Thr Leu Phe Leu Val Tyr
1415 1420 1425

Ser Asn Lys Cys Gln Thr Pro Leu Gly Met Ala Ser Gly His Ile
1430 1435 1440

Arg Asp Phe Gln Ile Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala
1445 1450 1455

Pro Lys Leu Ala Arg Leu His Tyr Ser Gly Ser Ile Asn Ala Trp
1460 1465 1470

Ser Thr Lys Glu Pro Phe Ser Trp Ile Lys Val Asp Leu Leu Ala
1475 1480 1485

Pro Met Ile Ile His Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys
1490 1495 1500

Phe Ser Ser Leu Tyr Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu
1505 1510 1515

Asp Gly Lys Lys Trp Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr

1520 1525 1530

Leu Met Val Phe Phe Gly Asn Val Asp Ser Ser Gly Ile Lys His
1535 1540 1545

Asn Ile Phe Asn Pro Pro Ile Ile Ala Arg Tyr Ile Arg Leu His
1550 1555 1560

Pro Thr His Tyr Ser Ile Arg Ser Thr Leu Arg Met Glu Leu Met
1565 1570 1575

Gly Cys Asp Leu Asn Ser Cys Ser Met Pro Leu Gly Met Glu Ser
1580 1585 1590

Lys Ala Ile Ser Asp Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr
1595 1600 1605

Asn Met Phe Ala Thr Trp Ser Pro Ser Lys Ala Arg Leu His Leu
1610 1615 1620

Gln Gly Arg Ser Asn Ala Trp Arg Pro Gln Val Asn Asn Pro Lys
1625 1630 1635

Glu Trp Leu Gln Val Asp Phe Gln Lys Thr Met Lys Val Thr Gly
1640 1645 1650

Val Thr Thr Gln Gly Val Lys Ser Leu Leu Thr Ser Met Tyr Val
1655 1660 1665

Lys Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly His Gln Trp Thr
1670 1675 1680

Leu Phe Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn Gln
1685 1690 1695

Asp Ser Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu
1700 1705 1710

Thr Arg Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile
1715 1720 1725

Ala Leu Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr
1730 1735 1740

Pro Leu Gly Arg Ile Val Gly Gly Gly Ala Ser Pro Gly Thr Ser
1745 1750 1755

Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr
1760 1765 1770

Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly

1775 1780 1785

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
1790 1795 1800

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
1805 1810 1815

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
1820 1825 1830

Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
1835 1840 1845

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
1850 1855 1860

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser
1865 1870 1875

Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
1880 1885 1890

Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
1895 1900 1905

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro
1910 1915 1920

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
1925 1930 1935

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
1940 1945 1950

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
1955 1960 1965

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Thr
1970 1975 1980

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly
1985 1990 1995

Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
2000 2005 2010

Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
2015 2020 2025

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly

2030 2035 2040

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser
2045 2050 2055

Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
2060 2065 2070

Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
2075 2080 2085

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
2090 2095 2100

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser
2105 2110 2115

Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
2120 2125 2130

Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
2135 2140 2145

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
2150 2155 2160

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
2165 2170 2175

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser
2180 2185 2190

Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
2195 2200 2205

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
2210 2215 2220

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
2225 2230 2235

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
2240 2245 2250

Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
2255 2260 2265

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
2270 2275 2280

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser

2285 2290 2295

Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
2300 2305 2310

Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
2315 2320 2325

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
2330 2335 2340

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
2345 2350 2355

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly
2360 2365 2370

Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
2375 2380 2385

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
2390 2395 2400

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
2405 2410 2415

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr
2420 2425 2430

Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
2435 2440 2445

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
2450 2455 2460

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
2465 2470 2475

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr
2480 2485 2490

Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly
2495 2500 2505

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
2510 2515 2520

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
2525 2530 2535

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly

2540 2545 2550

Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
2555 2560 2565

Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
2570 2575 2580

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly
2585 2590 2595

Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly
2600 2605 2610

Ser Pro
2615

<210> 1567
<211> 2242
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1567
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Leu Gln Val Arg Ile Val Gly Gly Gly Ala Pro Ser Pro Ser Ala Ser
130 135 140

```
Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
145                 150                 155                 160

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Pro Ser
                165                 170                 175

Gly Pro Gly Leu Gln Val Arg Ile Val Gly Gly Phe Pro Gly Gly Ser
            180                 185                 190

His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser
        195                 200                 205

Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val
    210                 215                 220

Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu
225                 230                 235                 240

Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu
                245                 250                 255

Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys
            260                 265                 270

Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro
        275                 280                 285

Lys Met His Thr Val Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu
    290                 295                 300

Ile Gly Cys His Arg Lys Ser Val Tyr Trp His Val Ile Gly Met Gly
305                 310                 315                 320

Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu Gly His Thr Phe Leu
                325                 330                 335

Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe
            340                 345                 350

Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe
        355                 360                 365

Cys His Ile Ser Ser His Gln His Asp Gly Met Glu Ala Tyr Val Lys
    370                 375                 380

Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu
385                 390                 395                 400

Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val
                405                 410                 415
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Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser
            420                 425                 430

Val Ala Lys Lys His Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu
        435                 440                 445

Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg
    450                 455                 460

Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg
465                 470                 475                 480

Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys
                485                 490                 495

Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu
            500                 505                 510

Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala
        515                 520                 525

Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro
    530                 535                 540

Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe
545                 550                 555                 560

Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val
                565                 570                 575

Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr
            580                 585                 590

Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly
        595                 600                 605

Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln
    610                 615                 620

Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu
625                 630                 635                 640

Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn
                645                 650                 655

Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile
            660                 665                 670

Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val
        675                 680                 685
```

```
Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln
    690                 695                 700

Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys
705                 710                 715                 720

Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr
                725                 730                 735

Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His
            740                 745                 750

Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser
        755                 760                 765

Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp
    770                 775                 780

Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser
785                 790                 795                 800

Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr
                805                 810                 815

Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr
            820                 825                 830

Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp
        835                 840                 845

Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe
    850                 855                 860

Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro
865                 870                 875                 880

His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys
                885                 890                 895

Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu
            900                 905                 910

Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile
        915                 920                 925

Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala
    930                 935                 940

Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp
945                 950                 955                 960
```

```
Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu
                965                 970                 975

Thr Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys
            980                 985                 990

Asp Glu Phe Asp Cys Lys Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu
        995                 1000                1005

Glu Lys  Asp Val His Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys
    1010                1015                1020

His Thr  Asn Thr Leu Asn Pro  Ala His Gly Arg Gln  Val Thr Val
    1025                1030                1035

Gln Glu  Phe Ala Leu Phe Phe  Thr Ile Phe Asp Glu  Thr Lys Ser
    1040                1045                1050

Trp Tyr  Phe Thr Glu Asn Met  Glu Arg Asn Cys Arg  Ala Pro Cys
    1055                1060                1065

Asn Ile  Gln Met Glu Asp Pro  Thr Phe Lys Glu Asn  Tyr Arg Phe
    1070                1075                1080

His Ala  Ile Asn Gly Tyr Ile  Met Asp Thr Leu Pro  Gly Leu Val
    1085                1090                1095

Met Ala  Gln Asp Gln Arg Ile  Arg Trp Tyr Leu Leu  Ser Met Gly
    1100                1105                1110

Ser Asn  Glu Asn Ile His Ser  Ile His Phe Ser Gly  His Val Phe
    1115                1120                1125

Thr Val  Arg Lys Lys Glu Glu  Tyr Lys Met Ala Leu  Tyr Asn Leu
    1130                1135                1140

Tyr Pro  Gly Val Phe Glu Thr  Val Glu Met Leu Pro  Ser Lys Ala
    1145                1150                1155

Gly Ile  Trp Arg Val Glu Cys  Leu Ile Gly Glu His  Leu His Ala
    1160                1165                1170

Gly Met  Ser Thr Leu Phe Leu  Val Tyr Ser Asn Lys  Cys Gln Thr
    1175                1180                1185

Pro Leu  Gly Met Ala Ser Gly  His Ile Arg Asp Phe  Gln Ile Thr
    1190                1195                1200

Ala Ser  Gly Gln Tyr Gly Leu  Gln Val Arg Ile Val  Gly Gly Ser
    1205                1210                1215
```

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Gly Thr  Ala Ser Ser Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1220                 1225                 1230

Thr Gly  Ser Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    1235                 1240                 1245

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser
    1250                 1255                 1260

Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Ser  Ser Pro Ser
    1265                 1270                 1275

Ala Ser  Thr Gly Thr Gly Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1280                 1285                 1290

Gly Thr  Gly Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    1295                 1300                 1305

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser
    1310                 1315                 1320

Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Ser  Ser Pro Ser
    1325                 1330                 1335

Ala Ser  Thr Gly Thr Gly Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1340                 1345                 1350

Gly Thr  Gly Pro Gly Ala Ser  Pro Gly Leu Gln Val  Arg Ile Val
    1355                 1360                 1365

Gly Gly  Gln Trp Ala Pro Lys  Leu Ala Arg Leu His  Tyr Ser Gly
    1370                 1375                 1380

Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro Phe Ser  Trp Ile Lys
    1385                 1390                 1395

Val Asp  Leu Leu Ala Pro Met  Ile Ile His Gly Ile  Lys Thr Gln
    1400                 1405                 1410

Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr Ile Ser  Gln Phe Ile
    1415                 1420                 1425

Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp Gln Thr  Tyr Arg Gly
    1430                 1435                 1440

Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe Gly Asn  Val Asp Ser
    1445                 1450                 1455

Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro Pro Ile  Ile Ala Arg
    1460                 1465                 1470
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Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser Ile Arg  Ser Thr Leu
    1475                1480                1485

Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn Ser Cys  Ser Met Pro
    1490                1495                1500

Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp Ala Gln  Ile Thr Ala
    1505                1510                1515

Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr Trp Ser  Pro Ser Lys
    1520                1525                1530

Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn Ala Trp  Arg Pro Gln
    1535                1540                1545

Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val Asp Phe  Gln Lys Thr
    1550                1555                1560

Met Lys  Val Thr Gly Val Thr  Thr Gln Gly Val Lys  Ser Leu Leu
    1565                1570                1575

Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile Ser Ser  Ser Gln Asp
    1580                1585                1590

Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn Gly Lys  Val Lys Val
    1595                1600                1605

Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro Val Val  Asn Ser Leu
    1610                1615                1620

Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg Ile His  Pro Gln Ser
    1625                1630                1635

Trp Val  His Gln Ile Ala Leu  Arg Met Glu Val Leu  Gly Cys Glu
    1640                1645                1650

Ala Gln  Asp Leu Tyr Leu Gln  Val Arg Ile Val Gly  Gly Pro Gly
    1655                1660                1665

Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser Pro Gly  Ser Ser Thr
    1670                1675                1680

Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser
    1685                1690                1695

Thr Gly  Thr Gly Pro Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr
    1700                1705                1710

Gly Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly
    1715                1720                1725
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Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser Pro Gly  Ala Ser Pro
    1730                1735                1740

Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1745                1750                1755

Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly
    1760                1765                1770

Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly
    1775                1780                1785

Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser Pro Gly  Ala Ser Pro
    1790                1795                1800

Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1805                1810                1815

Ser Thr  Gly Ser Pro Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr
    1820                1825                1830

Gly Pro  Gly Thr Pro Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly
    1835                1840                1845

Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser Pro Gly  Ala Ser Pro
    1850                1855                1860

Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1865                1870                1875

Ser Thr  Gly Ser Pro Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly
    1880                1885                1890

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly
    1895                1900                1905

Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser Pro Gly  Thr Pro Gly
    1910                1915                1920

Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser Ser Thr  Pro Ser Gly
    1925                1930                1935

Ala Thr  Gly Ser Pro Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly
    1940                1945                1950

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly
    1955                1960                1965

Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly Pro Gly  Ala Ser Pro
    1970                1975                1980
```

```
Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    1985                 1990                 1995

Ser Thr  Gly Ser Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
    2000                 2005                 2010

Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly
    2015                 2020                 2025

Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser Pro Gly  Ala Ser Pro
    2030                 2035                 2040

Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser
    2045                 2050                 2055

Ser Thr  Gly Ser Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
    2060                 2065                 2070

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly
    2075                 2080                 2085

Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser Pro Gly  Ser Ser Thr
    2090                 2095                 2100

Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Thr Pro Gly  Ser Gly Thr
    2105                 2110                 2115

Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly
    2120                 2125                 2130

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly
    2135                 2140                 2145

Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly Pro Gly  Ser Ser Pro
    2150                 2155                 2160

Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ala Ser Pro  Gly Thr Ser
    2165                 2170                 2175

Ser Thr  Gly Ser Pro Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser
    2180                 2185                 2190

Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly
    2195                 2200                 2205

Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly Pro Gly  Ser Ser Pro
    2210                 2215                 2220

Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ala Ser Pro  Gly Thr Ser
    2225                 2230                 2235
```

Ser Thr Gly Ser
2240

<210> 1568
<211> 3351
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1568
Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
35 40 45

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
50 55 60

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
65 70 75 80

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
85 90 95

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
100 105 110

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
115 120 125

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
130 135 140

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
145 150 155 160

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
165 170 175

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
180 185 190

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
195 200 205

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro

210 215 220

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
225 230 235 240

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
245 250 255

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
260 265 270

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
275 280 285

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
290 295 300

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
305 310 315 320

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
325 330 335

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
340 345 350

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
355 360 365

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
370 375 380

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
385 390 395 400

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
405 410 415

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
420 425 430

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
435 440 445

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
450 455 460

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
465 470 475 480

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr

485 490 495

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
500 505 510

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
515 520 525

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala
530 535 540

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
545 550 555 560

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
565 570 575

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
580 585 590

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
595 600 605

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
610 615 620

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
625 630 635 640

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
645 650 655

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
660 665 670

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
675 680 685

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
690 695 700

Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
705 710 715 720

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
725 730 735

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
740 745 750

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro

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        755                 760                 765

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
    770                 775                 780

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
785                 790                 795                 800

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
                805                 810                 815

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
            820                 825                 830

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
        835                 840                 845

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    850                 855                 860

Gly Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp
865                 870                 875                 880

Tyr Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro
                885                 890                 895

Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys
            900                 905                 910

Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys
        915                 920                 925

Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu
    930                 935                 940

Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro
945                 950                 955                 960

Val Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly
                965                 970                 975

Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys
            980                 985                 990

Val Phe Pro Gly Gly Ser His Thr  Tyr Val Trp Gln Val  Leu Lys Glu
        995                 1000                1005

Asn Gly  Pro Met Ala Ser Asp  Pro Leu Cys Leu Thr  Tyr Ser Tyr
    1010                1015                1020

Leu Ser  His Val Asp Leu Val  Lys Asp Leu Asn Ser  Gly Leu Ile
```

1025 1030 1035

Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys
1040 1045 1050

Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp
1055 1060 1065

Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
1070 1075 1080

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr
1085 1090 1095

Val Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys
1100 1105 1110

His Arg Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr
1115 1120 1125

Pro Glu Val His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val
1130 1135 1140

Arg Asn His Arg Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe
1145 1150 1155

Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly Gln Phe Leu Leu
1160 1165 1170

Phe Cys His Ile Ser Ser His Gln His Asp Gly Met Glu Ala Tyr
1175 1180 1185

Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg Met Lys
1190 1195 1200

Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp Ser
1205 1210 1215

Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
1220 1225 1230

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val
1235 1240 1245

His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu
1250 1255 1260

Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn
1265 1270 1275

Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe

```
    1280                1285                1290

Met Ala  Tyr Thr Asp Glu Thr  Phe Lys Thr Arg Glu  Ala Ile Gln
    1295                1300                1305

His Glu  Ser Gly Ile Leu Gly  Pro Leu Leu Tyr Gly  Glu Val Gly
    1310                1315                1320

Asp Thr  Leu Leu Ile Ile Phe  Lys Asn Gln Ala Ser  Arg Pro Tyr
    1325                1330                1335

Asn Ile  Tyr Pro His Gly Ile  Thr Asp Val Arg Pro  Leu Tyr Ser
    1340                1345                1350

Arg Arg  Leu Pro Lys Gly Val  Lys His Leu Lys Asp  Phe Pro Ile
    1355                1360                1365

Leu Pro  Gly Glu Ile Phe Lys  Tyr Lys Trp Thr Val  Thr Val Glu
    1370                1375                1380

Asp Gly  Pro Thr Lys Ser Asp  Pro Arg Cys Leu Thr  Arg Tyr Tyr
    1385                1390                1395

Ser Ser  Phe Val Asn Met Glu  Arg Asp Leu Ala Ser  Gly Leu Ile
    1400                1405                1410

Gly Pro  Leu Leu Ile Cys Tyr  Lys Glu Ser Val Asp  Gln Arg Gly
    1415                1420                1425

Asn Gln  Ile Met Ser Asp Lys  Arg Asn Val Ile Leu  Phe Ser Val
    1430                1435                1440

Phe Asp  Glu Asn Arg Ser Trp  Tyr Leu Thr Glu Asn  Ile Gln Arg
    1445                1450                1455

Phe Leu  Pro Asn Pro Ala Gly  Val Gln Leu Glu Asp  Pro Glu Phe
    1460                1465                1470

Gln Ala  Ser Asn Ile Met His  Ser Ile Asn Gly Tyr  Val Phe Asp
    1475                1480                1485

Ser Leu  Gln Leu Ser Val Cys  Leu His Glu Val Ala  Tyr Trp Tyr
    1490                1495                1500

Ile Leu  Ser Ile Gly Ala Gln  Thr Asp Phe Leu Ser  Val Phe Phe
    1505                1510                1515

Ser Gly  Tyr Thr Phe Lys His  Lys Met Val Tyr Glu  Asp Thr Leu
    1520                1525                1530

Thr Leu  Phe Pro Phe Ser Gly  Glu Thr Val Phe Met  Ser Met Glu
```

1535 1540 1545

Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
1550 1555 1560

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys
1565 1570 1575

Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala
1580 1585 1590

Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser
1595 1600 1605

Gln Asn Ser Arg His Pro Ser Thr Arg Gln Lys Gln Phe Asn Ala
1610 1615 1620

Thr Thr Ile Pro Glu Asn Asp Ile Glu Lys Thr Asp Pro Trp Phe
1625 1630 1635

Ala His Arg Thr Pro Met Pro Lys Ile Gln Asn Val Ser Ser Ser
1640 1645 1650

Asp Leu Leu Met Leu Leu Arg Gln Ser Pro Thr Pro His Gly Leu
1655 1660 1665

Ser Leu Ser Asp Leu Gln Glu Ala Lys Tyr Glu Thr Phe Ser Asp
1670 1675 1680

Asp Pro Ser Pro Gly Ala Ile Asp Ser Asn Asn Ser Leu Ser Glu
1685 1690 1695

Met Thr His Phe Arg Pro Gln Leu His His Ser Gly Asp Met Val
1700 1705 1710

Phe Thr Pro Glu Ser Gly Leu Gln Leu Arg Leu Asn Glu Lys Leu
1715 1720 1725

Gly Thr Thr Ala Ala Thr Glu Leu Lys Lys Leu Asp Phe Lys Val
1730 1735 1740

Ser Ser Thr Ser Asn Asn Leu Ile Ser Thr Ile Pro Ser Asp Asn
1745 1750 1755

Leu Ala Ala Gly Thr Asp Asn Thr Ser Ser Leu Gly Pro Pro Ser
1760 1765 1770

Met Pro Val His Tyr Asp Ser Gln Leu Asp Thr Thr Leu Phe Gly
1775 1780 1785

Lys Lys Ser Ser Pro Leu Thr Glu Ser Gly Gly Pro Leu Ser Leu

1790 1795 1800

Ser Glu Glu Asn Asn Asp Ser Lys Leu Leu Glu Ser Gly Leu Met
1805 1810 1815

Asn Ser Gln Glu Ser Ser Trp Gly Lys Asn Val Ser Ser Thr Glu
1820 1825 1830

Ser Gly Arg Leu Phe Lys Gly Lys Arg Ala His Gly Pro Ala Leu
1835 1840 1845

Leu Thr Lys Asp Asn Ala Leu Phe Lys Val Ser Ile Ser Leu Leu
1850 1855 1860

Lys Thr Asn Lys Thr Ser Asn Asn Ser Ala Thr Asn Arg Lys Thr
1865 1870 1875

His Ile Asp Gly Pro Ser Leu Leu Ile Glu Asn Ser Pro Ser Val
1880 1885 1890

Trp Gln Asn Ile Leu Glu Ser Asp Thr Glu Phe Lys Lys Val Thr
1895 1900 1905

Pro Leu Ile His Asp Arg Met Leu Met Asp Lys Asn Ala Thr Ala
1910 1915 1920

Leu Arg Leu Asn His Met Ser Asn Lys Thr Thr Ser Ser Lys Asn
1925 1930 1935

Met Glu Met Val Gln Gln Lys Lys Glu Gly Pro Ile Pro Pro Asp
1940 1945 1950

Ala Gln Asn Pro Asp Met Ser Phe Phe Lys Met Leu Phe Leu Pro
1955 1960 1965

Glu Ser Ala Arg Trp Ile Gln Arg Thr His Gly Lys Asn Ser Leu
1970 1975 1980

Asn Ser Gly Gln Gly Pro Ser Pro Lys Gln Leu Val Ser Leu Gly
1985 1990 1995

Pro Glu Lys Ser Val Glu Gly Gln Asn Phe Leu Ser Glu Lys Asn
2000 2005 2010

Lys Val Val Val Gly Lys Gly Glu Phe Thr Lys Asp Val Gly Leu
2015 2020 2025

Lys Glu Met Val Phe Pro Ser Ser Arg Asn Leu Phe Leu Thr Asn
2030 2035 2040

Leu Asp Asn Leu His Glu Asn Asn Thr His Asn Gln Glu Lys Lys

2045 2050 2055

Ile Gln Glu Glu Ile Glu Lys Lys Glu Thr Leu Ile Gln Glu Asn
2060 2065 2070

Val Val Leu Pro Gln Ile His Thr Val Thr Gly Thr Lys Asn Phe
2075 2080 2085

Met Lys Asn Leu Phe Leu Leu Ser Thr Arg Gln Asn Val Glu Gly
2090 2095 2100

Ser Tyr Asp Gly Ala Tyr Ala Pro Val Leu Gln Asp Phe Arg Ser
2105 2110 2115

Leu Asn Asp Ser Thr Asn Arg Thr Lys Lys His Thr Ala His Phe
2120 2125 2130

Ser Lys Lys Gly Glu Glu Glu Asn Leu Glu Gly Leu Gly Asn Gln
2135 2140 2145

Thr Lys Gln Ile Val Glu Lys Tyr Ala Cys Thr Thr Arg Ile Ser
2150 2155 2160

Pro Asn Thr Ser Gln Gln Asn Phe Val Thr Gln Arg Ser Lys Arg
2165 2170 2175

Ala Leu Lys Gln Phe Arg Leu Pro Leu Glu Glu Thr Glu Leu Glu
2180 2185 2190

Lys Arg Ile Ile Val Asp Asp Thr Ser Thr Gln Trp Ser Lys Asn
2195 2200 2205

Met Lys His Leu Thr Pro Ser Thr Leu Thr Gln Ile Asp Tyr Asn
2210 2215 2220

Glu Lys Glu Lys Gly Ala Ile Thr Gln Ser Pro Leu Ser Asp Cys
2225 2230 2235

Leu Thr Arg Ser His Ser Ile Pro Gln Ala Asn Arg Ser Pro Leu
2240 2245 2250

Pro Ile Ala Lys Val Ser Ser Phe Pro Ser Ile Arg Pro Ile Tyr
2255 2260 2265

Leu Thr Arg Val Leu Phe Gln Asp Asn Ser Ser His Leu Pro Ala
2270 2275 2280

Ala Ser Tyr Arg Lys Lys Asp Ser Gly Val Gln Glu Ser Ser His
2285 2290 2295

Phe Leu Gln Gly Ala Lys Lys Asn Asn Leu Ser Leu Ala Ile Leu

2300 2305 2310

Thr Leu Glu Met Thr Gly Asp Gln Arg Glu Val Gly Ser Leu Gly
2315 2320 2325

Thr Ser Ala Thr Asn Ser Val Thr Tyr Lys Lys Val Glu Asn Thr
2330 2335 2340

Val Leu Pro Lys Pro Asp Leu Pro Lys Thr Ser Gly Lys Val Glu
2345 2350 2355

Leu Leu Pro Lys Val His Ile Tyr Gln Lys Asp Leu Phe Pro Thr
2360 2365 2370

Glu Thr Ser Asn Gly Ser Pro Gly His Leu Asp Leu Val Glu Gly
2375 2380 2385

Ser Leu Leu Gln Gly Thr Glu Gly Ala Ile Lys Trp Asn Glu Ala
2390 2395 2400

Asn Arg Pro Gly Lys Val Pro Phe Leu Arg Val Ala Thr Glu Ser
2405 2410 2415

Ser Ala Lys Thr Pro Ser Lys Leu Leu Asp Pro Leu Ala Trp Asp
2420 2425 2430

Asn His Tyr Gly Thr Gln Ile Pro Lys Glu Glu Trp Lys Ser Gln
2435 2440 2445

Glu Lys Ser Pro Glu Lys Thr Ala Phe Lys Lys Lys Asp Thr Ile
2450 2455 2460

Leu Ser Leu Asn Ala Cys Glu Ser Asn His Ala Ile Ala Ala Ile
2465 2470 2475

Asn Glu Gly Gln Asn Lys Pro Glu Ile Glu Val Thr Trp Ala Lys
2480 2485 2490

Gln Gly Arg Thr Glu Arg Leu Cys Ser Gln Asn Pro Pro Val Leu
2495 2500 2505

Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp
2510 2515 2520

Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys
2525 2530 2535

Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
2540 2545 2550

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val

2555 2560 2565

Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu
2570 2575 2580

Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val
2585 2590 2595

Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
2600 2605 2610

Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile
2615 2620 2625

Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln
2630 2635 2640

Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu
2645 2650 2655

Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys
2660 2665 2670

Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His Met
2675 2680 2685

Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
2690 2695 2700

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly
2705 2710 2715

Pro Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly
2720 2725 2730

Arg Gln Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe
2735 2740 2745

Asp Glu Thr Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn
2750 2755 2760

Cys Arg Ala Pro Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys
2765 2770 2775

Glu Asn Tyr Arg Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr
2780 2785 2790

Leu Pro Gly Leu Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr
2795 2800 2805

Leu Leu Ser Met Gly Ser Asn Glu Asn Ile His Ser Ile His Phe

2810 2815 2820

Ser Gly His Val Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met
2825 2830 2835

Ala Leu Tyr Asn Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met
2840 2845 2850

Leu Pro Ser Lys Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly
2855 2860 2865

Glu His Leu His Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser
2870 2875 2880

Asn Lys Cys Gln Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg
2885 2890 2895

Asp Phe Gln Ile Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro
2900 2905 2910

Lys Leu Ala Arg Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser
2915 2920 2925

Thr Lys Glu Pro Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro
2930 2935 2940

Met Ile Ile His Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe
2945 2950 2955

Ser Ser Leu Tyr Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp
2960 2965 2970

Gly Lys Lys Trp Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu
2975 2980 2985

Met Val Phe Phe Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn
2990 2995 3000

Ile Phe Asn Pro Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro
3005 3010 3015

Thr His Tyr Ser Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly
3020 3025 3030

Cys Asp Leu Asn Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys
3035 3040 3045

Ala Ile Ser Asp Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn
3050 3055 3060

Met Phe Ala Thr Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln

3065 3070 3075

Gly Arg Ser Asn Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu
3080 3085 3090

Trp Leu Gln Val Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val
3095 3100 3105

Thr Thr Gln Gly Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys
3110 3115 3120

Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu
3125 3130 3135

Phe Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp
3140 3145 3150

Ser Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr
3155 3160 3165

Arg Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile Ala
3170 3175 3180

Leu Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly
3185 3190 3195

Leu Thr Pro Arg Ser Leu Leu Val Gly Gly Ser Glu Pro Ala Thr
3200 3205 3210

Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
3215 3220 3225

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
3230 3235 3240

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
3245 3250 3255

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr
3260 3265 3270

Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
3275 3280 3285

Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
3290 3295 3300

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
3305 3310 3315

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr

3320 3325 3330

Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
3335 3340 3345

Ser Ala Pro
3350

<210> 1569
<211> 1592
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1569
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

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His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460
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Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735
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Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Gly Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005
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Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1025                1030                1035

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1040                1045                1050

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1055                1060                1065

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1070                1075                1080

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1085                1090                1095

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1100                1105                1110

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1115                1120                1125

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1130                1135                1140

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1145                1150                1155

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1160                1165                1170

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1175                1180                1185

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1190                1195                1200

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1205                1210                1215

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1220                1225                1230

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1235                1240                1245

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1250                1255                1260
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Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1265                 1270                 1275

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1280                 1285                 1290

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1295                 1300                 1305

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1310                 1315                 1320

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1325                 1330                 1335

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1340                 1345                 1350

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1355                 1360                 1365

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1370                 1375                 1380

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1385                 1390                 1395

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1400                 1405                 1410

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1415                 1420                 1425

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Thr  Met Thr Arg
    1430                 1435                 1440

Ile Val  Gly Gly Gly Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu
    1445                 1450                 1455

Thr Pro  Gly Thr Ser Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly
    1460                 1465                 1470

Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr Pro Gly  Ser Pro Ala
    1475                 1480                 1485

Gly Ser  Pro Thr Ser Thr Glu  Glu Gly Thr Ser Thr  Glu Pro Ser
    1490                 1495                 1500

Glu Gly  Ser Ala Pro Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu
    1505                 1510                 1515
```

Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
1520 1525 1530

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr
1535 1540 1545

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
1550 1555 1560

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu
1565 1570 1575

Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
1580 1585 1590

<210> 1570
<211> 1584
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1570
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser

145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
340 345 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
355 360 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
370 375 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385 390 395 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
405 410 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr

420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
595 600 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala

690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
740 745 750

Gln Gly Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
755 760 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
770 775 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785 790 795 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
805 810 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
820 825 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
835 840 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
850 855 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865 870 875 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
885 890 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
900 905 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
915 920 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
930 935 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945 950 955 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln

965 970 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
980 985 990

Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
995 1000 1005

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1010 1015 1020

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1025 1030 1035

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1040 1045 1050

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1055 1060 1065

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1070 1075 1080

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1085 1090 1095

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1100 1105 1110

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1115 1120 1125

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1130 1135 1140

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg
1145 1150 1155

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1160 1165 1170

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1175 1180 1185

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1190 1195 1200

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1205 1210 1215

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe

1220 1225 1230

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1235 1240 1245

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1250 1255 1260

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1265 1270 1275

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1280 1285 1290

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1295 1300 1305

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1310 1315 1320

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1325 1330 1335

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1340 1345 1350

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1355 1360 1365

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1370 1375 1380

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1385 1390 1395

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1400 1405 1410

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1415 1420 1425

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Gly Gly Ser Glu
1430 1435 1440

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala
1445 1450 1455

Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
1460 1465 1470

Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu

1475 1480 1485

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu
1490 1495 1500

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr
1505 1510 1515

Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
1520 1525 1530

Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
1535 1540 1545

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
1550 1555 1560

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro
1565 1570 1575

Ser Glu Gly Ser Ala Pro
1580

<210> 1571
<211> 1591
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1571
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

```
Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380
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Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655
```

```
Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Gly Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925
```

```
Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1025                1030                1035

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1040                1045                1050

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1055                1060                1065

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1070                1075                1080

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1085                1090                1095

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1100                1105                1110

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1115                1120                1125

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1130                1135                1140

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1145                1150                1155

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1160                1165                1170

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1175                1180                1185
```

```
Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1190                1195                1200

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1205                1210                1215

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1220                1225                1230

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1235                1240                1245

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1250                1255                1260

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1265                1270                1275

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1280                1285                1290

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1295                1300                1305

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1310                1315                1320

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1325                1330                1335

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1340                1345                1350

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1355                1360                1365

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1370                1375                1380

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1385                1390                1395

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1400                1405                1410

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1415                1420                1425

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Gly Lys  Leu Thr Arg
    1430                1435                1440
```

```
Ala Glu  Thr Gly Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1445                1450                1455

Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser
    1460                1465                1470

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Ser  Pro Ala Gly
    1475                1480                1485

Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu
    1490                1495                1500

Gly Ser  Ala Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1505                1510                1515

Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Ser
    1520                1525                1530

Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro Gly Thr  Ser Thr Glu
    1535                1540                1545

Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser Glu Ser  Ala Thr Pro
    1550                1555                1560

Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr Ser Gly  Ser Glu Thr
    1565                1570                1575

Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly Ser Ala  Pro
    1580                1585                1590
```

<210> 1572
<211> 1592
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1572

```
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1               5                   10                  15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
```

65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225 230 235 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
245 250 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
260 265 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
275 280 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
290 295 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305 310 315 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
325 330 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp

```
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
```

610 615 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
675 680 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
690 695 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705 710 715 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
725 730 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
740 745 750

Gln Gly Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
755 760 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
770 775 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785 790 795 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
805 810 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
820 825 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
835 840 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
850 855 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865 870 875 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile

885 890 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
900 905 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
915 920 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
930 935 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945 950 955 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
965 970 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
980 985 990

Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys Arg Ala Pro
995 1000 1005

Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg
1010 1015 1020

Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu
1025 1030 1035

Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met
1040 1045 1050

Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser Gly His Val
1055 1060 1065

Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn
1070 1075 1080

Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu Pro Ser Lys
1085 1090 1095

Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu His Leu His
1100 1105 1110

Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln
1115 1120 1125

Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp Phe Gln Ile
1130 1135 1140

Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg

1145 1150 1155

Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro
1160 1165 1170

Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His
1175 1180 1185

Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr
1190 1195 1200

Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp
1205 1210 1215

Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met Val Phe Phe
1220 1225 1230

Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile Phe Asn Pro
1235 1240 1245

Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr His Tyr Ser
1250 1255 1260

Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys Asp Leu Asn
1265 1270 1275

Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala Ile Ser Asp
1280 1285 1290

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1295 1300 1305

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1310 1315 1320

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1325 1330 1335

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1340 1345 1350

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1355 1360 1365

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1370 1375 1380

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1385 1390 1395

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg

1400 1405 1410

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1415 1420 1425

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Leu Thr Pro Arg
1430 1435 1440

Ser Leu Leu Val Gly Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu
1445 1450 1455

Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
1460 1465 1470

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
1475 1480 1485

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
1490 1495 1500

Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu
1505 1510 1515

Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
1520 1525 1530

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr
1535 1540 1545

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
1550 1555 1560

Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu
1565 1570 1575

Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
1580 1585 1590

<210> 1573
<211> 2457
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1573
Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
20 25 30

```
Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
        35                  40                  45

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
    50                  55                  60

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
65                  70                  75                  80

Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
                85                  90                  95

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
            100                 105                 110

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
        115                 120                 125

Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    130                 135                 140

Gly Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp
145                 150                 155                 160

Tyr Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro
                165                 170                 175

Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys
            180                 185                 190

Lys Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys
        195                 200                 205

Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu
    210                 215                 220

Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro
225                 230                 235                 240

Val Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly
                245                 250                 255

Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys
            260                 265                 270

Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu
        275                 280                 285

Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu
    290                 295                 300
```

```
Ser His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala
305                 310                 315                 320

Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr
                325                 330                 335

Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser
            340                 345                 350

Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala
        355                 360                 365

Ser Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn
    370                 375                 380

Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp
385                 390                 395                 400

His Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu
                405                 410                 415

Glu Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu
            420                 425                 430

Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu
        435                 440                 445

Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly
    450                 455                 460

Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu
465                 470                 475                 480

Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr
                485                 490                 495

Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser
            500                 505                 510

Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val
        515                 520                 525

His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val
    530                 535                 540

Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly
545                 550                 555                 560

Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr
                565                 570                 575
```

```
Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly
            580                 585                 590

Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile
        595                 600                 605

Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly
    610                 615                 620

Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val
625                 630                 635                 640

Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr
                645                 650                 655

Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg
            660                 665                 670

Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu
        675                 680                 685

Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val
    690                 695                 700

Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu
705                 710                 715                 720

Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile
                725                 730                 735

Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu
            740                 745                 750

Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp
        755                 760                 765

Ser Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile
    770                 775                 780

Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly
785                 790                 795                 800

Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe
                805                 810                 815

Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu
            820                 825                 830

Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr
        835                 840                 845
```

```
Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr
    850                 855                 860

Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn
865                 870                 875                 880

Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg
                885                 890                 895

His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu
            900                 905                 910

Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
        915                 920                 925

Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys
    930                 935                 940

Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr
945                 950                 955                 960

Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly
                965                 970                 975

Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly
            980                 985                 990

Ser Phe Thr Gln Pro Leu Tyr Arg  Gly Glu Leu Asn Glu  His Leu Gly
        995                 1000                1005

Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val Glu Asp  Asn Ile Met
    1010                1015                1020

Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro Tyr Ser  Phe Tyr Ser
    1025                1030                1035

Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg Gln Gly  Ala Glu Pro
    1040                1045                1050

Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr Lys Thr  Tyr Phe Trp
    1055                1060                1065

Lys Val  Gln His His Met Ala  Pro Thr Lys Asp Glu  Phe Asp Cys
    1070                1075                1080

Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp Leu Glu  Lys Asp Val
    1085                1090                1095

His Ser  Gly Leu Ile Gly Pro  Leu Leu Val Cys His  Thr Asn Thr
    1100                1105                1110
```

```
Leu Asn  Pro Ala His Gly Arg  Gln Val Thr Val Gln  Glu Phe Ala
    1115                1120                1125

Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys Ser Trp  Tyr Phe Thr
    1130                1135                1140

Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro Cys Asn  Ile Gln Met
    1145                1150                1155

Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg Phe His  Ala Ile Asn
    1160                1165                1170

Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu Val Met  Ala Gln Asp
    1175                1180                1185

Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met Gly Ser  Asn Glu Asn
    1190                1195                1200

Ile His  Ser Ile His Phe Ser  Gly His Val Phe Thr  Val Arg Lys
    1205                1210                1215

Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn Leu Tyr  Pro Gly Val
    1220                1225                1230

Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys Ala Gly  Ile Trp Arg
    1235                1240                1245

Val Glu  Cys Leu Ile Gly Glu  His Leu His Ala Gly  Met Ser Thr
    1250                1255                1260

Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln Thr Pro  Leu Gly Met
    1265                1270                1275

Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile Thr Ala  Ser Gly Gln
    1280                1285                1290

Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg Leu His  Tyr Ser Gly
    1295                1300                1305

Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro Phe Ser  Trp Ile Lys
    1310                1315                1320

Val Asp  Leu Leu Ala Pro Met  Ile Ile His Gly Ile  Lys Thr Gln
    1325                1330                1335

Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr Ile Ser  Gln Phe Ile
    1340                1345                1350

Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp Gln Thr  Tyr Arg Gly
    1355                1360                1365
```

```
Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe Gly Asn  Val Asp Ser
    1370                1375                1380

Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro Pro Ile  Ile Ala Arg
    1385                1390                1395

Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser Ile Arg  Ser Thr Leu
    1400                1405                1410

Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn Ser Cys  Ser Met Pro
    1415                1420                1425

Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp Ala Gln  Ile Thr Ala
    1430                1435                1440

Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr Trp Ser  Pro Ser Lys
    1445                1450                1455

Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn Ala Trp  Arg Pro Gln
    1460                1465                1470

Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val Asp Phe  Gln Lys Thr
    1475                1480                1485

Met Lys  Val Thr Gly Val Thr  Thr Gln Gly Val Lys  Ser Leu Leu
    1490                1495                1500

Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile Ser Ser  Ser Gln Asp
    1505                1510                1515

Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn Gly Lys  Val Lys Val
    1520                1525                1530

Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro Val Val  Asn Ser Leu
    1535                1540                1545

Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg Ile His  Pro Gln Ser
    1550                1555                1560

Trp Val  His Gln Ile Ala Leu  Arg Met Glu Val Leu  Gly Cys Glu
    1565                1570                1575

Ala Gln  Asp Leu Tyr Gly Ala  Pro Leu Gly Leu Arg  Leu Arg Gly
    1580                1585                1590

Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser
    1595                1600                1605

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro
    1610                1615                1620
```

```
Ser Glu  Gly Ser Ala Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1625                 1630                 1635

Thr Glu  Glu Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1640                 1645                 1650

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1655                 1660                 1665

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
    1670                 1675                 1680

Ser Gly  Ser Glu Thr Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1685                 1690                 1695

Glu Thr  Pro Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1700                 1705                 1710

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1715                 1720                 1725

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Thr Glu Pro
    1730                 1735                 1740

Ser Glu  Gly Ser Ala Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1745                 1750                 1755

Thr Glu  Glu Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1760                 1765                 1770

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1775                 1780                 1785

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro
    1790                 1795                 1800

Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1805                 1810                 1815

Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro
    1820                 1825                 1830

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1835                 1840                 1845

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala
    1850                 1855                 1860

Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1865                 1870                 1875
```

```
Ser Gly  Pro Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1880                1885                1890

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    1895                1900                1905

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1910                1915                1920

Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1925                1930                1935

Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1940                1945                1950

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1955                1960                1965

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Thr Glu Pro
    1970                1975                1980

Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1985                1990                1995

Ser Ala  Pro Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    2000                2005                2010

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    2015                2020                2025

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
    2030                2035                2040

Ser Gly  Ser Glu Thr Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    2045                2050                2055

Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro
    2060                2065                2070

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    2075                2080                2085

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala
    2090                2095                2100

Thr Pro  Glu Ser Gly Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    2105                2110                2115

Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    2120                2125                2130
```

```
Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser
    2135                2140                2145

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro
    2150                2155                2160

Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    2165                2170                2175

Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro
    2180                2185                2190

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    2195                2200                2205

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    2210                2215                2220

Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    2225                2230                2235

Ser Ala  Pro Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    2240                2245                2250

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    2255                2260                2265

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    2270                2275                2280

Thr Pro  Glu Ser Gly Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    2285                2290                2295

Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    2300                2305                2310

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    2315                2320                2325

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Thr Ser  Glu Ser Ala
    2330                2335                2340

Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    2345                2350                2355

Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro
    2360                2365                2370

Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Pro
    2375                2380                2385
```

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro
    2390   2395   2400

Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
    2405   2410   2415

Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
    2420   2425   2430

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
    2435   2440   2445

Thr Glu Pro Ser Glu Gly Ser Ala Pro
    2450   2455

<210> 1574
<211> 2457
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
      polypeptide

<400> 1574
Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
1   5   10   15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
            20   25   30

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
        35   40   45

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
    50   55   60

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
65   70   75   80

Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
                85   90   95

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
            100   105   110

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
        115   120   125

Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    130   135   140

Gly Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp

```
145                 150                 155                 160

Tyr Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro
                165                 170                 175

Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys
            180                 185                 190

Lys Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys
        195                 200                 205

Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu
    210                 215                 220

Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro
225                 230                 235                 240

Val Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly
                245                 250                 255

Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys
            260                 265                 270

Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu
        275                 280                 285

Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu
    290                 295                 300

Ser His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala
305                 310                 315                 320

Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr
                325                 330                 335

Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser
            340                 345                 350

Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala
        355                 360                 365

Ser Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn
    370                 375                 380

Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp
385                 390                 395                 400

His Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu
                405                 410                 415

Glu Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu
```

```
            420                 425                 430

Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu
        435                 440                 445

Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly
    450                 455                 460

Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu
465                 470                 475                 480

Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr
                485                 490                 495

Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser
            500                 505                 510

Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val
        515                 520                 525

His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val
    530                 535                 540

Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly
545                 550                 555                 560

Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr
                565                 570                 575

Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly
            580                 585                 590

Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile
        595                 600                 605

Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly
    610                 615                 620

Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val
625                 630                 635                 640

Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr
                645                 650                 655

Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg
            660                 665                 670

Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu
        675                 680                 685

Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val
```

690 695 700

Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu
705 710 715 720

Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile
725 730 735

Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu
740 745 750

Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp
755 760 765

Ser Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile
770 775 780

Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly
785 790 795 800

Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe
805 810 815

Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu
820 825 830

Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr
835 840 845

Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr
850 855 860

Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn
865 870 875 880

Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg
885 890 895

His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu
900 905 910

Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
915 920 925

Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys
930 935 940

Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr
945 950 955 960

Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly

965 970 975

Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly
980 985 990

Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly
995 1000 1005

Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met
1010 1015 1020

Val Thr Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser
1025 1030 1035

Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro
1040 1045 1050

Arg Lys Asn Phe Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp
1055 1060 1065

Lys Val Gln His His Met Ala Pro Thr Lys Asp Glu Phe Asp Cys
1070 1075 1080

Lys Ala Trp Ala Tyr Phe Ser Asp Val Asp Leu Glu Lys Asp Val
1085 1090 1095

His Ser Gly Leu Ile Gly Pro Leu Leu Val Cys His Thr Asn Thr
1100 1105 1110

Leu Asn Pro Ala His Gly Arg Gln Val Thr Val Gln Glu Phe Ala
1115 1120 1125

Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys Ser Trp Tyr Phe Thr
1130 1135 1140

Glu Asn Met Glu Arg Asn Cys Arg Ala Pro Cys Asn Ile Gln Met
1145 1150 1155

Glu Asp Pro Thr Phe Lys Glu Asn Tyr Arg Phe His Ala Ile Asn
1160 1165 1170

Gly Tyr Ile Met Asp Thr Leu Pro Gly Leu Val Met Ala Gln Asp
1175 1180 1185

Gln Arg Ile Arg Trp Tyr Leu Leu Ser Met Gly Ser Asn Glu Asn
1190 1195 1200

Ile His Ser Ile His Phe Ser Gly His Val Phe Thr Val Arg Lys
1205 1210 1215

Lys Glu Glu Tyr Lys Met Ala Leu Tyr Asn Leu Tyr Pro Gly Val

1220 1225 1230

Phe Glu Thr Val Glu Met Leu Pro Ser Lys Ala Gly Ile Trp Arg
1235 1240 1245

Val Glu Cys Leu Ile Gly Glu His Leu His Ala Gly Met Ser Thr
1250 1255 1260

Leu Phe Leu Val Tyr Ser Asn Lys Cys Gln Thr Pro Leu Gly Met
1265 1270 1275

Ala Ser Gly His Ile Arg Asp Phe Gln Ile Thr Ala Ser Gly Gln
1280 1285 1290

Tyr Gly Gln Trp Ala Pro Lys Leu Ala Arg Leu His Tyr Ser Gly
1295 1300 1305

Ser Ile Asn Ala Trp Ser Thr Lys Glu Pro Phe Ser Trp Ile Lys
1310 1315 1320

Val Asp Leu Leu Ala Pro Met Ile Ile His Gly Ile Lys Thr Gln
1325 1330 1335

Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr Ile Ser Gln Phe Ile
1340 1345 1350

Ile Met Tyr Ser Leu Asp Gly Lys Lys Trp Gln Thr Tyr Arg Gly
1355 1360 1365

Asn Ser Thr Gly Thr Leu Met Val Phe Phe Gly Asn Val Asp Ser
1370 1375 1380

Ser Gly Ile Lys His Asn Ile Phe Asn Pro Pro Ile Ile Ala Arg
1385 1390 1395

Tyr Ile Arg Leu His Pro Thr His Tyr Ser Ile Arg Ser Thr Leu
1400 1405 1410

Arg Met Glu Leu Met Gly Cys Asp Leu Asn Ser Cys Ser Met Pro
1415 1420 1425

Leu Gly Met Glu Ser Lys Ala Ile Ser Asp Ala Gln Ile Thr Ala
1430 1435 1440

Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp Ser Pro Ser Lys
1445 1450 1455

Ala Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg Pro Gln
1460 1465 1470

Val Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln Lys Thr

1475 1480 1485

Met Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser Leu Leu
1490 1495 1500

Thr Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser Gln Asp
1505 1510 1515

Gly His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val Lys Val
1520 1525 1530

Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn Ser Leu
1535 1540 1545

Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro Gln Ser
1550 1555 1560

Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly Cys Glu
1565 1570 1575

Ala Gln Asp Leu Tyr Gly Thr Met Thr Arg Ile Val Gly Gly Gly
1580 1585 1590

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
1595 1600 1605

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro
1610 1615 1620

Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
1625 1630 1635

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
1640 1645 1650

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
1655 1660 1665

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr
1670 1675 1680

Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
1685 1690 1695

Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
1700 1705 1710

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
1715 1720 1725

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro

1730 1735 1740

Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
1745 1750 1755

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
1760 1765 1770

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
1775 1780 1785

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro
1790 1795 1800

Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
1805 1810 1815

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
1820 1825 1830

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
1835 1840 1845

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala
1850 1855 1860

Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
1865 1870 1875

Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
1880 1885 1890

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
1895 1900 1905

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala
1910 1915 1920

Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
1925 1930 1935

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
1940 1945 1950

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
1955 1960 1965

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro
1970 1975 1980

Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly

1985 1990 1995

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
2000 2005 2010

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
2015 2020 2025

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr
2030 2035 2040

Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
2045 2050 2055

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
2060 2065 2070

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
2075 2080 2085

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala
2090 2095 2100

Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
2105 2110 2115

Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
2120 2125 2130

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
2135 2140 2145

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro
2150 2155 2160

Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
2165 2170 2175

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
2180 2185 2190

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
2195 2200 2205

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala
2210 2215 2220

Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
2225 2230 2235

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu

2240 2245 2250

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu
2255 2260 2265

Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala
2270 2275 2280

Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
2285 2290 2295

Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
2300 2305 2310

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
2315 2320 2325

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala
2330 2335 2340

Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
2345 2350 2355

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
2360 2365 2370

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro
2375 2380 2385

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro
2390 2395 2400

Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
2405 2410 2415

Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
2420 2425 2430

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
2435 2440 2445

Thr Glu Pro Ser Glu Gly Ser Ala Pro
2450 2455

<210> 1575
<211> 2168
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1575

```
Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
1               5                  10                  15

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
            20                  25                  30

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr
        35                  40                  45

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
    50                  55                  60

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
65                  70                  75                  80

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly
                85                  90                  95

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
            100                 105                 110

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
        115                 120                 125

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
    130                 135                 140

Gly Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp
145                 150                 155                 160

Tyr Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro
                165                 170                 175

Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys
            180                 185                 190

Lys Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys
        195                 200                 205

Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu
    210                 215                 220

Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro
225                 230                 235                 240

Val Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly
                245                 250                 255

Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys
            260                 265                 270
```

```
Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu
        275                 280                 285

Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu
    290                 295                 300

Ser His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala
305                 310                 315                 320

Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr
                325                 330                 335

Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser
            340                 345                 350

Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala
        355                 360                 365

Ser Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn
    370                 375                 380

Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp
385                 390                 395                 400

His Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu
                405                 410                 415

Glu Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu
            420                 425                 430

Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu
        435                 440                 445

Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly
    450                 455                 460

Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu
465                 470                 475                 480

Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr
                485                 490                 495

Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser
            500                 505                 510

Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val
        515                 520                 525

His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val
    530                 535                 540
```

```
Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly
545                 550                 555                 560

Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr
                565                 570                 575

Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly
            580                 585                 590

Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile
        595                 600                 605

Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly
    610                 615                 620

Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val
625                 630                 635                 640

Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr
                645                 650                 655

Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg
            660                 665                 670

Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu
        675                 680                 685

Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val
    690                 695                 700

Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu
705                 710                 715                 720

Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile
                725                 730                 735

Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu
            740                 745                 750

Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp
        755                 760                 765

Ser Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile
    770                 775                 780

Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly
785                 790                 795                 800

Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe
                805                 810                 815
```

```
Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu
            820                 825                 830

Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr
        835                 840                 845

Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr
    850                 855                 860

Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn
865                 870                 875                 880

Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg
                885                 890                 895

His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu
            900                 905                 910

Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
        915                 920                 925

Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys
    930                 935                 940

Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr
945                 950                 955                 960

Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly
                965                 970                 975

Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly
            980                 985                 990

Ser Phe Thr Gln Pro Leu Tyr Arg  Gly Glu Leu Asn Glu  His Leu Gly
        995                 1000                1005

Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val Glu Asp  Asn Ile Met
    1010                1015                1020

Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro Tyr Ser  Phe Tyr Ser
    1025                1030                1035

Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg Gln Gly  Ala Glu Pro
    1040                1045                1050

Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr Lys Thr  Tyr Phe Trp
    1055                1060                1065

Lys Val  Gln His His Met Ala  Pro Thr Lys Asp Glu  Phe Asp Cys
    1070                1075                1080
```

```
Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp Leu Glu  Lys Asp Val
    1085                1090                1095

His Ser  Gly Leu Ile Gly Pro  Leu Leu Val Cys His  Thr Asn Thr
    1100                1105                1110

Leu Asn  Pro Ala His Gly Arg  Gln Val Thr Val Gln  Glu Phe Ala
    1115                1120                1125

Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys Ser Trp  Tyr Phe Thr
    1130                1135                1140

Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro Cys Asn  Ile Gln Met
    1145                1150                1155

Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg Phe His  Ala Ile Asn
    1160                1165                1170

Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu Val Met  Ala Gln Asp
    1175                1180                1185

Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met Gly Ser  Asn Glu Asn
    1190                1195                1200

Ile His  Ser Ile His Phe Ser  Gly His Val Phe Thr  Val Arg Lys
    1205                1210                1215

Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn Leu Tyr  Pro Gly Val
    1220                1225                1230

Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys Ala Gly  Ile Trp Arg
    1235                1240                1245

Val Glu  Cys Leu Ile Gly Glu  His Leu His Ala Gly  Met Ser Thr
    1250                1255                1260

Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln Thr Pro  Leu Gly Met
    1265                1270                1275

Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile Thr Ala  Ser Gly Gln
    1280                1285                1290

Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg Leu His  Tyr Ser Gly
    1295                1300                1305

Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro Phe Ser  Trp Ile Lys
    1310                1315                1320

Val Asp  Leu Leu Ala Pro Met  Ile Ile His Gly Ile  Lys Thr Gln
    1325                1330                1335
```

```
Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr Ile Ser  Gln Phe Ile
    1340                1345                1350

Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp Gln Thr  Tyr Arg Gly
    1355                1360                1365

Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe Gly Asn  Val Asp Ser
    1370                1375                1380

Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro Pro Ile  Ile Ala Arg
    1385                1390                1395

Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser Ile Arg  Ser Thr Leu
    1400                1405                1410

Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn Ser Cys  Ser Met Pro
    1415                1420                1425

Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp Ala Gln  Ile Thr Ala
    1430                1435                1440

Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr Trp Ser  Pro Ser Lys
    1445                1450                1455

Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn Ala Trp  Arg Pro Gln
    1460                1465                1470

Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val Asp Phe  Gln Lys Thr
    1475                1480                1485

Met Lys  Val Thr Gly Val Thr  Thr Gln Gly Val Lys  Ser Leu Leu
    1490                1495                1500

Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile Ser Ser  Ser Gln Asp
    1505                1510                1515

Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn Gly Lys  Val Lys Val
    1520                1525                1530

Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro Val Val  Asn Ser Leu
    1535                1540                1545

Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg Ile His  Pro Gln Ser
    1550                1555                1560

Trp Val  His Gln Ile Ala Leu  Arg Met Glu Val Leu  Gly Cys Glu
    1565                1570                1575

Ala Gln  Asp Leu Tyr Gly Lys  Leu Thr Arg Ala Glu  Thr Gly Pro
    1580                1585                1590
```

```
Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser Ser Pro  Gly Ser Ser
    1595                 1600                 1605

Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala
    1610                 1615                 1620

Ser Thr  Gly Thr Gly Pro Gly  Ser Ser Pro Ser Ala  Ser Thr Gly
    1625                 1630                 1635

Thr Gly  Pro Gly Ser Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro
    1640                 1645                 1650

Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Gly Ala Ser
    1655                 1660                 1665

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    1670                 1675                 1680

Ser Ser  Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
    1685                 1690                 1695

Gly Ser  Pro Gly Thr Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro
    1700                 1705                 1710

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1715                 1720                 1725

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    1730                 1735                 1740

Ser Ser  Thr Gly Ser Pro Gly  Ser Ser Pro Ser Ala  Ser Thr Gly
    1745                 1750                 1755

Thr Gly  Pro Gly Thr Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro
    1760                 1765                 1770

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1775                 1780                 1785

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    1790                 1795                 1800

Ser Ser  Thr Gly Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    1805                 1810                 1815

Gly Ser  Pro Gly Ser Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro
    1820                 1825                 1830

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Thr Pro
    1835                 1840                 1845
```

```
Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly Ser Ser  Thr Pro Ser
    1850                1855                1860

Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    1865                1870                1875

Gly Ser  Pro Gly Ser Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro
    1880                1885                1890

Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr Gly Pro  Gly Ala Ser
    1895                1900                1905

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    1910                1915                1920

Ser Ser  Thr Gly Ser Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    1925                1930                1935

Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1940                1945                1950

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1955                1960                1965

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    1970                1975                1980

Ser Ser  Thr Gly Ser Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    1985                1990                1995

Ser Ser  Pro Gly Ser Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro
    2000                2005                2010

Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser Ser Pro  Gly Ser Ser
    2015                2020                2025

Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Thr Pro  Gly Ser Gly
    2030                2035                2040

Thr Ala  Ser Ser Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    2045                2050                2055

Gly Ser  Pro Gly Ser Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro
    2060                2065                2070

Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr Gly Pro  Gly Ser Ser
    2075                2080                2085

Pro Ser  Ala Ser Thr Gly Thr  Gly Pro Gly Ala Ser  Pro Gly Thr
    2090                2095                2100
```

Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser
2105 2110 2115

Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
2120 2125 2130

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser
2135 2140 2145

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr
2150 2155 2160

Ser Ser Thr Gly Ser
2165

<210> 1576
<211> 2456
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1576
Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
20 25 30

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
35 40 45

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
50 55 60

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
65 70 75 80

Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
85 90 95

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
100 105 110

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
115 120 125

Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
130 135 140

Gly Lys Leu Thr Arg Ala Glu Thr Gly Ala Thr Arg Arg Tyr Tyr Leu

145 150 155 160

Gly Ala Val Glu Leu Ser Trp Asp Tyr Met Gln Ser Asp Leu Gly Glu
165 170 175

Leu Pro Val Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro
180 185 190

Phe Asn Thr Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr
195 200 205

Asp His Leu Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu
210 215 220

Leu Gly Pro Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr
225 230 235 240

Leu Lys Asn Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val
245 250 255

Ser Tyr Trp Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser
260 265 270

Gln Arg Glu Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr
275 280 285

Tyr Val Trp Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro
290 295 300

Leu Cys Leu Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp
305 310 315 320

Leu Asn Ser Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser
325 330 335

Leu Ala Lys Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe
340 345 350

Ala Val Phe Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser
355 360 365

Leu Met Gln Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met
370 375 380

His Thr Val Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly
385 390 395 400

Cys His Arg Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr
405 410 415

Pro Glu Val His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg

420 425 430

Asn His Arg Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr
435 440 445

Ala Gln Thr Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His
450 455 460

Ile Ser Ser His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp
465 470 475 480

Ser Cys Pro Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala
485 490 495

Glu Asp Tyr Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg
500 505 510

Phe Asp Asp Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala
515 520 525

Lys Lys His Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu
530 535 540

Asp Trp Asp Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr
545 550 555 560

Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr
565 570 575

Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg
580 585 590

Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly
595 600 605

Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg
610 615 620

Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr
625 630 635 640

Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile
645 650 655

Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp
660 665 670

Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser
675 680 685

Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu

690 695 700

Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met
705 710 715 720

Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg
725 730 735

Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala
740 745 750

Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His
755 760 765

Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu
770 775 780

His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp
785 790 795 800

Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val
805 810 815

Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe
820 825 830

Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser
835 840 845

Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys
850 855 860

Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser
865 870 875 880

Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser
885 890 895

Gln Asn Pro Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr
900 905 910

Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser
915 920 925

Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn
930 935 940

Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala
945 950 955 960

Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val

965 970 975

Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val
980 985 990

Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg
995 1000 1005

Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
1010 1015 1020

Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala
1025 1030 1035

Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu
1040 1045 1050

Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro
1055 1060 1065

Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala
1070 1075 1080

Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe Ser
1085 1090 1095

Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
1100 1105 1110

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg
1115 1120 1125

Gln Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp
1130 1135 1140

Glu Thr Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn Cys
1145 1150 1155

Arg Ala Pro Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys Glu
1160 1165 1170

Asn Tyr Arg Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr Leu
1175 1180 1185

Pro Gly Leu Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr Leu
1190 1195 1200

Leu Ser Met Gly Ser Asn Glu Asn Ile His Ser Ile His Phe Ser
1205 1210 1215

Gly His Val Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met Ala

1220 1225 1230

Leu Tyr Asn Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met Leu
1235 1240 1245

Pro Ser Lys Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly Glu
1250 1255 1260

His Leu His Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser Asn
1265 1270 1275

Lys Cys Gln Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg Asp
1280 1285 1290

Phe Gln Ile Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro Lys
1295 1300 1305

Leu Ala Arg Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr
1310 1315 1320

Lys Glu Pro Phe Ser Trp Ile Lys Val Asp Leu Leu Ala Pro Met
1325 1330 1335

Ile Ile His Gly Ile Lys Thr Gln Gly Ala Arg Gln Lys Phe Ser
1340 1345 1350

Ser Leu Tyr Ile Ser Gln Phe Ile Ile Met Tyr Ser Leu Asp Gly
1355 1360 1365

Lys Lys Trp Gln Thr Tyr Arg Gly Asn Ser Thr Gly Thr Leu Met
1370 1375 1380

Val Phe Phe Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn Ile
1385 1390 1395

Phe Asn Pro Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro Thr
1400 1405 1410

His Tyr Ser Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly Cys
1415 1420 1425

Asp Leu Asn Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys Ala
1430 1435 1440

Ile Ser Asp Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met
1445 1450 1455

Phe Ala Thr Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly
1460 1465 1470

Arg Ser Asn Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp

1475 1480 1485

Leu Gln Val Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr
1490 1495 1500

Thr Gln Gly Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu
1505 1510 1515

Phe Leu Ile Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe
1520 1525 1530

Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser
1535 1540 1545

Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg
1550 1555 1560

Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu
1565 1570 1575

Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly Gly
1580 1585 1590

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
1595 1600 1605

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser
1610 1615 1620

Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr
1625 1630 1635

Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
1640 1645 1650

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
1655 1660 1665

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
1670 1675 1680

Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu
1685 1690 1695

Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
1700 1705 1710

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
1715 1720 1725

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser

1730 1735 1740

Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr
1745 1750 1755

Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
1760 1765 1770

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
1775 1780 1785

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser
1790 1795 1800

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
1805 1810 1815

Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
1820 1825 1830

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
1835 1840 1845

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
1850 1855 1860

Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
1865 1870 1875

Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
1880 1885 1890

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
1895 1900 1905

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
1910 1915 1920

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
1925 1930 1935

Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
1940 1945 1950

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
1955 1960 1965

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser
1970 1975 1980

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser

1985 1990 1995

Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
2000 2005 2010

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
2015 2020 2025

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser
2030 2035 2040

Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
2045 2050 2055

Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
2060 2065 2070

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
2075 2080 2085

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr
2090 2095 2100

Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr
2105 2110 2115

Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
2120 2125 2130

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
2135 2140 2145

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser
2150 2155 2160

Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
2165 2170 2175

Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
2180 2185 2190

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
2195 2200 2205

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
2210 2215 2220

Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
2225 2230 2235

Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly

2240 2245 2250

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
2255 2260 2265

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr
2270 2275 2280

Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr
2285 2290 2295

Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
2300 2305 2310

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
2315 2320 2325

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr
2330 2335 2340

Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
2345 2350 2355

Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
2360 2365 2370

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
2375 2380 2385

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
2390 2395 2400

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
2405 2410 2415

Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
2420 2425 2430

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
2435 2440 2445

Glu Pro Ser Glu Gly Ser Ala Pro
2450 2455

<210> 1577
<211> 2457
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1577

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Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
1               5                   10                  15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
            20                  25                  30

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
        35                  40                  45

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
    50                  55                  60

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
65                  70                  75                  80

Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
                85                  90                  95

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
            100                 105                 110

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
        115                 120                 125

Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    130                 135                 140

Gly Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp
145                 150                 155                 160

Tyr Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro
                165                 170                 175

Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys
            180                 185                 190

Lys Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys
        195                 200                 205

Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu
    210                 215                 220

Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro
225                 230                 235                 240

Val Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly
                245                 250                 255

Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys
            260                 265                 270
```

```
Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu
        275                 280                 285

Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu
    290                 295                 300

Ser His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala
305                 310                 315                 320

Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr
                325                 330                 335

Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser
            340                 345                 350

Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala
        355                 360                 365

Ser Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn
    370                 375                 380

Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp
385                 390                 395                 400

His Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu
                405                 410                 415

Glu Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu
            420                 425                 430

Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu
        435                 440                 445

Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly
    450                 455                 460

Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu
465                 470                 475                 480

Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr
                485                 490                 495

Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser
            500                 505                 510

Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val
        515                 520                 525

His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val
    530                 535                 540
```

```
Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly
545                 550                 555                 560

Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr
                565                 570                 575

Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly
            580                 585                 590

Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile
        595                 600                 605

Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly
    610                 615                 620

Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val
625                 630                 635                 640

Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr
                645                 650                 655

Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg
            660                 665                 670

Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu
        675                 680                 685

Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val
    690                 695                 700

Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu
705                 710                 715                 720

Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile
                725                 730                 735

Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu
            740                 745                 750

Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp
        755                 760                 765

Ser Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile
    770                 775                 780

Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly
785                 790                 795                 800

Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe
                805                 810                 815
```

```
Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu
            820                 825                 830

Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr
        835                 840                 845

Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr
    850                 855                 860

Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn
865                 870                 875                 880

Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg
                885                 890                 895

His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu
            900                 905                 910

Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe
        915                 920                 925

Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys
    930                 935                 940

Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr
945                 950                 955                 960

Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly
                965                 970                 975

Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly
            980                 985                 990

Ser Phe Thr Gln Pro Leu Tyr Arg  Gly Glu Leu Asn Glu  His Leu Gly
        995                 1000                1005

Leu Leu  Gly Pro Tyr Ile Arg  Ala Glu Val Glu Asp  Asn Ile Met
    1010                1015                1020

Val Thr  Phe Arg Asn Gln Ala  Ser Arg Pro Tyr Ser  Phe Tyr Ser
    1025                1030                1035

Ser Leu  Ile Ser Tyr Glu Glu  Asp Gln Arg Gln Gly  Ala Glu Pro
    1040                1045                1050

Arg Lys  Asn Phe Val Lys Pro  Asn Glu Thr Lys Thr  Tyr Phe Trp
    1055                1060                1065

Lys Val  Gln His His Met Ala  Pro Thr Lys Asp Glu  Phe Asp Cys
    1070                1075                1080
```

```
Lys Ala  Trp Ala Tyr Phe Ser  Asp Val Asp Leu Glu  Lys Asp Val
    1085                1090                1095

His Ser  Gly Leu Ile Gly Pro  Leu Leu Val Cys His  Thr Asn Thr
    1100                1105                1110

Leu Asn  Pro Ala His Gly Arg  Gln Val Thr Val Gln  Glu Phe Ala
    1115                1120                1125

Leu Phe  Phe Thr Ile Phe Asp  Glu Thr Lys Ser Trp  Tyr Phe Thr
    1130                1135                1140

Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro Cys Asn  Ile Gln Met
    1145                1150                1155

Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg Phe His  Ala Ile Asn
    1160                1165                1170

Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu Val Met  Ala Gln Asp
    1175                1180                1185

Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met Gly Ser  Asn Glu Asn
    1190                1195                1200

Ile His  Ser Ile His Phe Ser  Gly His Val Phe Thr  Val Arg Lys
    1205                1210                1215

Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn Leu Tyr  Pro Gly Val
    1220                1225                1230

Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys Ala Gly  Ile Trp Arg
    1235                1240                1245

Val Glu  Cys Leu Ile Gly Glu  His Leu His Ala Gly  Met Ser Thr
    1250                1255                1260

Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln Thr Pro  Leu Gly Met
    1265                1270                1275

Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile Thr Ala  Ser Gly Gln
    1280                1285                1290

Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg Leu His  Tyr Ser Gly
    1295                1300                1305

Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro Phe Ser  Trp Ile Lys
    1310                1315                1320

Val Asp  Leu Leu Ala Pro Met  Ile Ile His Gly Ile  Lys Thr Gln
    1325                1330                1335
```

```
Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr Ile Ser  Gln Phe Ile
    1340                1345                1350

Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp Gln Thr  Tyr Arg Gly
    1355                1360                1365

Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe Gly Asn  Val Asp Ser
    1370                1375                1380

Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro Pro Ile  Ile Ala Arg
    1385                1390                1395

Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser Ile Arg  Ser Thr Leu
    1400                1405                1410

Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn Ser Cys  Ser Met Pro
    1415                1420                1425

Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp Ala Gln  Ile Thr Ala
    1430                1435                1440

Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr Trp Ser  Pro Ser Lys
    1445                1450                1455

Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn Ala Trp  Arg Pro Gln
    1460                1465                1470

Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val Asp Phe  Gln Lys Thr
    1475                1480                1485

Met Lys  Val Thr Gly Val Thr  Thr Gln Gly Val Lys  Ser Leu Leu
    1490                1495                1500

Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile Ser Ser  Ser Gln Asp
    1505                1510                1515

Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn Gly Lys  Val Lys Val
    1520                1525                1530

Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro Val Val  Asn Ser Leu
    1535                1540                1545

Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg Ile His  Pro Gln Ser
    1550                1555                1560

Trp Val  His Gln Ile Ala Leu  Arg Met Glu Val Leu  Gly Cys Glu
    1565                1570                1575

Ala Gln  Asp Leu Tyr Gly Leu  Thr Pro Arg Ser Leu  Leu Val Gly
    1580                1585                1590
```

```
Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser
    1595                 1600                 1605

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro
    1610                 1615                 1620

Ser Glu  Gly Ser Ala Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1625                 1630                 1635

Thr Glu  Glu Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1640                 1645                 1650

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1655                 1660                 1665

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
    1670                 1675                 1680

Ser Gly  Ser Glu Thr Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1685                 1690                 1695

Glu Thr  Pro Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1700                 1705                 1710

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1715                 1720                 1725

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Thr Glu Pro
    1730                 1735                 1740

Ser Glu  Gly Ser Ala Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1745                 1750                 1755

Thr Glu  Glu Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1760                 1765                 1770

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1775                 1780                 1785

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro
    1790                 1795                 1800

Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1805                 1810                 1815

Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro
    1820                 1825                 1830

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1835                 1840                 1845
```

```
Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala
    1850                 1855                 1860

Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1865                 1870                 1875

Ser Gly  Pro Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1880                 1885                 1890

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    1895                 1900                 1905

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1910                 1915                 1920

Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1925                 1930                 1935

Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1940                 1945                 1950

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1955                 1960                 1965

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Thr Glu Pro
    1970                 1975                 1980

Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1985                 1990                 1995

Ser Ala  Pro Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    2000                 2005                 2010

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    2015                 2020                 2025

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Glu  Pro Ala Thr
    2030                 2035                 2040

Ser Gly  Ser Glu Thr Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    2045                 2050                 2055

Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro
    2060                 2065                 2070

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    2075                 2080                 2085

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala
    2090                 2095                 2100
```

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Thr Pro  Glu Ser Gly Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    2105                 2110                 2115

Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    2120                 2125                 2130

Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser
    2135                 2140                 2145

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro
    2150                 2155                 2160

Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    2165                 2170                 2175

Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro
    2180                 2185                 2190

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    2195                 2200                 2205

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    2210                 2215                 2220

Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    2225                 2230                 2235

Ser Ala  Pro Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    2240                 2245                 2250

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    2255                 2260                 2265

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    2270                 2275                 2280

Thr Pro  Glu Ser Gly Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    2285                 2290                 2295

Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    2300                 2305                 2310

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    2315                 2320                 2325

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Thr Ser  Glu Ser Ala
    2330                 2335                 2340

Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    2345                 2350                 2355
```

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
2360 2365 2370

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro
2375 2380 2385

Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro
2390 2395 2400

Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
2405 2410 2415

Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
2420 2425 2430

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
2435 2440 2445

Thr Glu Pro Ser Glu Gly Ser Ala Pro
2450 2455

<210> 1578
<211> 3351
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1578
Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
35 40 45

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
50 55 60

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
65 70 75 80

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
85 90 95

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
100 105 110

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro

115 120 125

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
130 135 140

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
145 150 155 160

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
165 170 175

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
180 185 190

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
195 200 205

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
210 215 220

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
225 230 235 240

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
245 250 255

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
260 265 270

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
275 280 285

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
290 295 300

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
305 310 315 320

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
325 330 335

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
340 345 350

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
355 360 365

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
370 375 380

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr

385 390 395 400

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
405 410 415

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
420 425 430

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
435 440 445

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
450 455 460

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
465 470 475 480

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
485 490 495

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
500 505 510

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
515 520 525

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala
530 535 540

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
545 550 555 560

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
565 570 575

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
580 585 590

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
595 600 605

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
610 615 620

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
625 630 635 640

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
645 650 655

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro

660 665 670

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
675 680 685

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
690 695 700

Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
705 710 715 720

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
725 730 735

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
740 745 750

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
755 760 765

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
770 775 780

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
785 790 795 800

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
805 810 815

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
820 825 830

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
835 840 845

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
850 855 860

Gly Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp
865 870 875 880

Tyr Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro
885 890 895

Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys
900 905 910

Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys
915 920 925

Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu

930 935 940

Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro
945 950 955 960

Val Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly
965 970 975

Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys
980 985 990

Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu
995 1000 1005

Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr
1010 1015 1020

Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile
1025 1030 1035

Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys
1040 1045 1050

Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp
1055 1060 1065

Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln
1070 1075 1080

Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr
1085 1090 1095

Val Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys
1100 1105 1110

His Arg Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr
1115 1120 1125

Pro Glu Val His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val
1130 1135 1140

Arg Asn His Arg Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe
1145 1150 1155

Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly Gln Phe Leu Leu
1160 1165 1170

Phe Cys His Ile Ser Ser His Gln His Asp Gly Met Glu Ala Tyr
1175 1180 1185

Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg Met Lys

1190 1195 1200

Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp Ser
1205 1210 1215

Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
1220 1225 1230

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val
1235 1240 1245

His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu
1250 1255 1260

Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn
1265 1270 1275

Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe
1280 1285 1290

Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln
1295 1300 1305

His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly
1310 1315 1320

Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr
1325 1330 1335

Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser
1340 1345 1350

Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile
1355 1360 1365

Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu
1370 1375 1380

Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr
1385 1390 1395

Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile
1400 1405 1410

Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly
1415 1420 1425

Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val
1430 1435 1440

Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg

1445 1450 1455

Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
1460 1465 1470

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp
1475 1480 1485

Ser Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr
1490 1495 1500

Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe
1505 1510 1515

Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu
1520 1525 1530

Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu
1535 1540 1545

Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg
1550 1555 1560

Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys
1565 1570 1575

Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala
1580 1585 1590

Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser
1595 1600 1605

Gln Asn Ser Arg His Pro Ser Thr Arg Gln Lys Gln Phe Asn Ala
1610 1615 1620

Thr Thr Ile Pro Glu Asn Asp Ile Glu Lys Thr Asp Pro Trp Phe
1625 1630 1635

Ala His Arg Thr Pro Met Pro Lys Ile Gln Asn Val Ser Ser Ser
1640 1645 1650

Asp Leu Leu Met Leu Leu Arg Gln Ser Pro Thr Pro His Gly Leu
1655 1660 1665

Ser Leu Ser Asp Leu Gln Glu Ala Lys Tyr Glu Thr Phe Ser Asp
1670 1675 1680

Asp Pro Ser Pro Gly Ala Ile Asp Ser Asn Asn Ser Leu Ser Glu
1685 1690 1695

Met Thr His Phe Arg Pro Gln Leu His His Ser Gly Asp Met Val

1700 1705 1710

Phe Thr Pro Glu Ser Gly Leu Gln Leu Arg Leu Asn Glu Lys Leu
1715 1720 1725

Gly Thr Thr Ala Ala Thr Glu Leu Lys Lys Leu Asp Phe Lys Val
1730 1735 1740

Ser Ser Thr Ser Asn Asn Leu Ile Ser Thr Ile Pro Ser Asp Asn
1745 1750 1755

Leu Ala Ala Gly Thr Asp Asn Thr Ser Ser Leu Gly Pro Pro Ser
1760 1765 1770

Met Pro Val His Tyr Asp Ser Gln Leu Asp Thr Thr Leu Phe Gly
1775 1780 1785

Lys Lys Ser Ser Pro Leu Thr Glu Ser Gly Gly Pro Leu Ser Leu
1790 1795 1800

Ser Glu Glu Asn Asn Asp Ser Lys Leu Leu Glu Ser Gly Leu Met
1805 1810 1815

Asn Ser Gln Glu Ser Ser Trp Gly Lys Asn Val Ser Ser Thr Glu
1820 1825 1830

Ser Gly Arg Leu Phe Lys Gly Lys Arg Ala His Gly Pro Ala Leu
1835 1840 1845

Leu Thr Lys Asp Asn Ala Leu Phe Lys Val Ser Ile Ser Leu Leu
1850 1855 1860

Lys Thr Asn Lys Thr Ser Asn Asn Ser Ala Thr Asn Arg Lys Thr
1865 1870 1875

His Ile Asp Gly Pro Ser Leu Leu Ile Glu Asn Ser Pro Ser Val
1880 1885 1890

Trp Gln Asn Ile Leu Glu Ser Asp Thr Glu Phe Lys Lys Val Thr
1895 1900 1905

Pro Leu Ile His Asp Arg Met Leu Met Asp Lys Asn Ala Thr Ala
1910 1915 1920

Leu Arg Leu Asn His Met Ser Asn Lys Thr Thr Ser Ser Lys Asn
1925 1930 1935

Met Glu Met Val Gln Gln Lys Lys Glu Gly Pro Ile Pro Pro Asp
1940 1945 1950

Ala Gln Asn Pro Asp Met Ser Phe Phe Lys Met Leu Phe Leu Pro

1955 1960 1965

Glu Ser Ala Arg Trp Ile Gln Arg Thr His Gly Lys Asn Ser Leu
1970 1975 1980

Asn Ser Gly Gln Gly Pro Ser Pro Lys Gln Leu Val Ser Leu Gly
1985 1990 1995

Pro Glu Lys Ser Val Glu Gly Gln Asn Phe Leu Ser Glu Lys Asn
2000 2005 2010

Lys Val Val Val Gly Lys Gly Glu Phe Thr Lys Asp Val Gly Leu
2015 2020 2025

Lys Glu Met Val Phe Pro Ser Ser Arg Asn Leu Phe Leu Thr Asn
2030 2035 2040

Leu Asp Asn Leu His Glu Asn Asn Thr His Asn Gln Glu Lys Lys
2045 2050 2055

Ile Gln Glu Glu Ile Glu Lys Lys Glu Thr Leu Ile Gln Glu Asn
2060 2065 2070

Val Val Leu Pro Gln Ile His Thr Val Thr Gly Thr Lys Asn Phe
2075 2080 2085

Met Lys Asn Leu Phe Leu Leu Ser Thr Arg Gln Asn Val Glu Gly
2090 2095 2100

Ser Tyr Asp Gly Ala Tyr Ala Pro Val Leu Gln Asp Phe Arg Ser
2105 2110 2115

Leu Asn Asp Ser Thr Asn Arg Thr Lys Lys His Thr Ala His Phe
2120 2125 2130

Ser Lys Lys Gly Glu Glu Glu Asn Leu Glu Gly Leu Gly Asn Gln
2135 2140 2145

Thr Lys Gln Ile Val Glu Lys Tyr Ala Cys Thr Thr Arg Ile Ser
2150 2155 2160

Pro Asn Thr Ser Gln Gln Asn Phe Val Thr Gln Arg Ser Lys Arg
2165 2170 2175

Ala Leu Lys Gln Phe Arg Leu Pro Leu Glu Glu Thr Glu Leu Glu
2180 2185 2190

Lys Arg Ile Ile Val Asp Asp Thr Ser Thr Gln Trp Ser Lys Asn
2195 2200 2205

Met Lys His Leu Thr Pro Ser Thr Leu Thr Gln Ile Asp Tyr Asn

2210 2215 2220

Glu Lys Glu Lys Gly Ala Ile Thr Gln Ser Pro Leu Ser Asp Cys
2225 2230 2235

Leu Thr Arg Ser His Ser Ile Pro Gln Ala Asn Arg Ser Pro Leu
2240 2245 2250

Pro Ile Ala Lys Val Ser Ser Phe Pro Ser Ile Arg Pro Ile Tyr
2255 2260 2265

Leu Thr Arg Val Leu Phe Gln Asp Asn Ser Ser His Leu Pro Ala
2270 2275 2280

Ala Ser Tyr Arg Lys Lys Asp Ser Gly Val Gln Glu Ser Ser His
2285 2290 2295

Phe Leu Gln Gly Ala Lys Lys Asn Asn Leu Ser Leu Ala Ile Leu
2300 2305 2310

Thr Leu Glu Met Thr Gly Asp Gln Arg Glu Val Gly Ser Leu Gly
2315 2320 2325

Thr Ser Ala Thr Asn Ser Val Thr Tyr Lys Lys Val Glu Asn Thr
2330 2335 2340

Val Leu Pro Lys Pro Asp Leu Pro Lys Thr Ser Gly Lys Val Glu
2345 2350 2355

Leu Leu Pro Lys Val His Ile Tyr Gln Lys Asp Leu Phe Pro Thr
2360 2365 2370

Glu Thr Ser Asn Gly Ser Pro Gly His Leu Asp Leu Val Glu Gly
2375 2380 2385

Ser Leu Leu Gln Gly Thr Glu Gly Ala Ile Lys Trp Asn Glu Ala
2390 2395 2400

Asn Arg Pro Gly Lys Val Pro Phe Leu Arg Val Ala Thr Glu Ser
2405 2410 2415

Ser Ala Lys Thr Pro Ser Lys Leu Leu Asp Pro Leu Ala Trp Asp
2420 2425 2430

Asn His Tyr Gly Thr Gln Ile Pro Lys Glu Glu Trp Lys Ser Gln
2435 2440 2445

Glu Lys Ser Pro Glu Lys Thr Ala Phe Lys Lys Lys Asp Thr Ile
2450 2455 2460

Leu Ser Leu Asn Ala Cys Glu Ser Asn His Ala Ile Ala Ala Ile

2465 2470 2475

Asn Glu Gly Gln Asn Lys Pro Glu Ile Glu Val Thr Trp Ala Lys
2480 2485 2490

Gln Gly Arg Thr Glu Arg Leu Cys Ser Gln Asn Pro Pro Val Leu
2495 2500 2505

Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp
2510 2515 2520

Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys
2525 2530 2535

Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
2540 2545 2550

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val
2555 2560 2565

Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu
2570 2575 2580

Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val
2585 2590 2595

Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
2600 2605 2610

Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile
2615 2620 2625

Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln
2630 2635 2640

Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu
2645 2650 2655

Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys
2660 2665 2670

Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His Met
2675 2680 2685

Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
2690 2695 2700

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly
2705 2710 2715

Pro Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly

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    2720                2725                2730

Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu Phe Phe  Thr Ile Phe
    2735                2740                2745

Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu Asn Met  Glu Arg Asn
    2750                2755                2760

Cys Arg  Ala Pro Cys Asn Ile  Gln Met Glu Asp Pro  Thr Phe Lys
    2765                2770                2775

Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly Tyr Ile  Met Asp Thr
    2780                2785                2790

Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln Arg Ile  Arg Trp Tyr
    2795                2800                2805

Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile His Ser  Ile His Phe
    2810                2815                2820

Ser Gly  His Val Phe Thr Val  Arg Lys Lys Glu Glu  Tyr Lys Met
    2825                2830                2835

Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe Glu Thr  Val Glu Met
    2840                2845                2850

Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val Glu Cys  Leu Ile Gly
    2855                2860                2865

Glu His  Leu His Ala Gly Met  Ser Thr Leu Phe Leu  Val Tyr Ser
    2870                2875                2880

Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala Ser Gly  His Ile Arg
    2885                2890                2895

Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr Gly Gln  Trp Ala Pro
    2900                2905                2910

Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser Ile Asn  Ala Trp Ser
    2915                2920                2925

Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val Asp Leu  Leu Ala Pro
    2930                2935                2940

Met Ile  Ile His Gly Ile Lys  Thr Gln Gly Ala Arg  Gln Lys Phe
    2945                2950                2955

Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile Met Tyr  Ser Leu Asp
    2960                2965                2970

Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn Ser Thr  Gly Thr Leu
```

2975 2980 2985

Met Val Phe Phe Gly Asn Val Asp Ser Ser Gly Ile Lys His Asn
2990 2995 3000

Ile Phe Asn Pro Pro Ile Ile Ala Arg Tyr Ile Arg Leu His Pro
3005 3010 3015

Thr His Tyr Ser Ile Arg Ser Thr Leu Arg Met Glu Leu Met Gly
3020 3025 3030

Cys Asp Leu Asn Ser Cys Ser Met Pro Leu Gly Met Glu Ser Lys
3035 3040 3045

Ala Ile Ser Asp Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn
3050 3055 3060

Met Phe Ala Thr Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln
3065 3070 3075

Gly Arg Ser Asn Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu
3080 3085 3090

Trp Leu Gln Val Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val
3095 3100 3105

Thr Thr Gln Gly Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys
3110 3115 3120

Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu
3125 3130 3135

Phe Phe Gln Asn Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp
3140 3145 3150

Ser Phe Thr Pro Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr
3155 3160 3165

Arg Tyr Leu Arg Ile His Pro Gln Ser Trp Val His Gln Ile Ala
3170 3175 3180

Leu Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr Gly
3185 3190 3195

Ala Pro Leu Gly Leu Arg Leu Arg Gly Gly Ser Glu Pro Ala Thr
3200 3205 3210

Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
3215 3220 3225

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro

3230 3235 3240

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
3245 3250 3255

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr
3260 3265 3270

Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
3275 3280 3285

Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
3290 3295 3300

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser
3305 3310 3315

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr
3320 3325 3330

Ser Gly Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
3335 3340 3345

Ser Ala Pro
3350

<210> 1579
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<212> PRT
<213> Artificial Sequence

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<222> (2924)..(2926)
<223> Any amino acid

<220>
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<222> (2937)..(2940)
<223> Any amino acid

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1 5 10 15

Ser Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser
20 25 30

Ser Thr Ala Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro
35 40 45

```
Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser
    50                  55                  60

Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro Ser Gly Glu Ser
65                  70                  75                  80

Ser Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
                85                  90                  95

Gly Ser Thr Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro
            100                 105                 110

Ser Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser
        115                 120                 125

Ser Thr Ala Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro
    130                 135                 140

Gly Thr Met Thr Arg Ile Val Gly Gly Ala Thr Arg Arg Tyr Tyr Leu
145                 150                 155                 160

Gly Ala Val Glu Leu Ser Trp Asp Tyr Met Gln Ser Asp Leu Gly Glu
                165                 170                 175

Leu Pro Val Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro
            180                 185                 190

Phe Asn Thr Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr
        195                 200                 205

Asp His Leu Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu
    210                 215                 220

Leu Gly Pro Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr
225                 230                 235                 240

Leu Lys Asn Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val
                245                 250                 255

Ser Tyr Trp Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser
            260                 265                 270

Gln Arg Glu Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr
        275                 280                 285

Tyr Val Trp Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro
    290                 295                 300

Leu Cys Leu Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp
305                 310                 315                 320
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Leu Asn Ser Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser
                325                 330                 335

Leu Ala Lys Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe
            340                 345                 350

Ala Val Phe Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser
        355                 360                 365

Leu Met Gln Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met
    370                 375                 380

His Thr Val Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly
385                 390                 395                 400

Cys His Arg Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr
                405                 410                 415

Pro Glu Val His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg
            420                 425                 430

Asn His Arg Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr
        435                 440                 445

Ala Gln Thr Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His
    450                 455                 460

Ile Ser Ser His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp
465                 470                 475                 480

Ser Cys Pro Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala
                485                 490                 495

Glu Asp Tyr Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg
            500                 505                 510

Phe Asp Asp Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala
        515                 520                 525

Lys Lys His Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu
    530                 535                 540

Asp Trp Asp Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr
545                 550                 555                 560

Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr
                565                 570                 575

Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg
            580                 585                 590
```

```
Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly
        595                 600                 605

Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg
    610                 615                 620

Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr
625                 630                 635                 640

Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile
                645                 650                 655

Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp
            660                 665                 670

Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser
        675                 680                 685

Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu
    690                 695                 700

Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met
705                 710                 715                 720

Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg
                725                 730                 735

Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala
            740                 745                 750

Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His
        755                 760                 765

Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu
    770                 775                 780

His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp
785                 790                 795                 800

Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val
                805                 810                 815

Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe
            820                 825                 830

Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser
        835                 840                 845

Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys
    850                 855                 860
```

```
Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser
865                 870                 875                 880

Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser
                885                 890                 895

Gln Asn Ser Arg His Pro Ser Thr Arg Gln Lys Gln Phe Asn Ala Thr
            900                 905                 910

Thr Ile Pro Glu Asn Asp Ile Glu Lys Thr Asp Pro Trp Phe Ala His
        915                 920                 925

Arg Thr Pro Met Pro Lys Ile Gln Asn Val Ser Ser Ser Asp Leu Leu
    930                 935                 940

Met Leu Leu Arg Gln Ser Pro Thr Pro His Gly Leu Ser Leu Ser Asp
945                 950                 955                 960

Leu Gln Glu Ala Lys Tyr Glu Thr Phe Ser Asp Asp Pro Ser Pro Gly
                965                 970                 975

Ala Ile Asp Ser Asn Asn Ser Leu Ser Glu Met Thr His Phe Arg Pro
            980                 985                 990

Gln Leu His His Ser Gly Asp Met  Val Phe Thr Pro Glu  Ser Gly Leu
        995                 1000                1005

Gln Leu  Arg Leu Asn Glu Lys  Leu Gly Thr Thr Ala  Ala Thr Glu
    1010                1015                1020

Leu Lys  Lys Leu Asp Phe Lys  Val Ser Ser Thr Ser  Asn Asn Leu
    1025                1030                1035

Ile Ser  Thr Ile Pro Ser Asp  Asn Leu Ala Ala Gly  Thr Asp Asn
    1040                1045                1050

Thr Ser  Ser Leu Gly Pro Pro  Ser Met Pro Val His  Tyr Asp Ser
    1055                1060                1065

Gln Leu  Asp Thr Thr Leu Phe  Gly Lys Lys Ser Ser  Pro Leu Thr
    1070                1075                1080

Glu Ser  Gly Gly Pro Leu Ser  Leu Ser Glu Glu Asn  Asn Asp Ser
    1085                1090                1095

Lys Leu  Leu Glu Ser Gly Leu  Met Asn Ser Gln Glu  Ser Ser Trp
    1100                1105                1110

Gly Lys  Asn Val Ser Ser Thr  Glu Ser Gly Arg Leu  Phe Lys Gly
    1115                1120                1125
```

```
Lys Arg  Ala His Gly Pro Ala  Leu Leu Thr Lys Asp  Asn Ala Leu
    1130                1135                1140

Phe Lys  Val Ser Ile Ser Leu  Leu Lys Thr Asn Lys  Thr Ser Asn
    1145                1150                1155

Asn Ser  Ala Thr Asn Arg Lys  Thr His Ile Asp Gly  Pro Ser Leu
    1160                1165                1170

Leu Ile  Glu Asn Ser Pro Ser  Val Trp Gln Asn Ile  Leu Glu Ser
    1175                1180                1185

Asp Thr  Glu Phe Lys Lys Val  Thr Pro Leu Ile His  Asp Arg Met
    1190                1195                1200

Leu Met  Asp Lys Asn Ala Thr  Ala Leu Arg Leu Asn  His Met Ser
    1205                1210                1215

Asn Lys  Thr Thr Ser Ser Lys  Asn Met Glu Met Val  Gln Gln Lys
    1220                1225                1230

Lys Glu  Gly Pro Ile Pro Pro  Asp Ala Gln Asn Pro  Asp Met Ser
    1235                1240                1245

Phe Phe  Lys Met Leu Phe Leu  Pro Glu Ser Ala Arg  Trp Ile Gln
    1250                1255                1260

Arg Thr  His Gly Lys Asn Ser  Leu Asn Ser Gly Gln  Gly Pro Ser
    1265                1270                1275

Pro Lys  Gln Leu Val Ser Leu  Gly Pro Glu Lys Ser  Val Glu Gly
    1280                1285                1290

Gln Asn  Phe Leu Ser Glu Lys  Asn Lys Val Val Val  Gly Lys Gly
    1295                1300                1305

Glu Phe  Thr Lys Asp Val Gly  Leu Lys Glu Met Val  Phe Pro Ser
    1310                1315                1320

Ser Arg  Asn Leu Phe Leu Thr  Asn Leu Asp Asn Leu  His Glu Asn
    1325                1330                1335

Asn Thr  His Asn Gln Glu Lys  Lys Ile Gln Glu Glu  Ile Glu Lys
    1340                1345                1350

Lys Glu  Thr Leu Ile Gln Glu  Asn Val Val Leu Pro  Gln Ile His
    1355                1360                1365

Thr Val  Thr Gly Thr Lys Asn  Phe Met Lys Asn Leu  Phe Leu Leu
    1370                1375                1380
```

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Ser Thr  Arg Gln Asn Val Glu  Gly Ser Tyr Asp Gly  Ala Tyr Ala
    1385                1390                1395

Pro Val  Leu Gln Asp Phe Arg  Ser Leu Asn Asp Ser  Thr Asn Arg
    1400                1405                1410

Thr Lys  Lys His Thr Ala His  Phe Ser Lys Lys Gly  Glu Glu Glu
    1415                1420                1425

Asn Leu  Glu Gly Leu Gly Asn  Gln Thr Lys Gln Ile  Val Glu Lys
    1430                1435                1440

Tyr Ala  Cys Thr Thr Arg Ile  Ser Pro Asn Thr Ser  Gln Gln Asn
    1445                1450                1455

Phe Val  Thr Gln Arg Ser Lys  Arg Ala Leu Lys Gln  Phe Arg Leu
    1460                1465                1470

Pro Leu  Glu Glu Thr Glu Leu  Glu Lys Arg Ile Ile  Val Asp Asp
    1475                1480                1485

Thr Ser  Thr Gln Trp Ser Lys  Asn Met Lys His Leu  Thr Pro Ser
    1490                1495                1500

Thr Leu  Thr Gln Ile Asp Tyr  Asn Glu Lys Glu Lys  Gly Ala Ile
    1505                1510                1515

Thr Gln  Ser Pro Leu Ser Asp  Cys Leu Thr Arg Ser  His Ser Ile
    1520                1525                1530

Pro Gln  Ala Asn Arg Ser Pro  Leu Pro Ile Ala Lys  Val Ser Ser
    1535                1540                1545

Phe Pro  Ser Ile Arg Pro Ile  Tyr Leu Thr Arg Val  Leu Phe Gln
    1550                1555                1560

Asp Asn  Ser Ser His Leu Pro  Ala Ala Ser Tyr Arg  Lys Lys Asp
    1565                1570                1575

Ser Gly  Val Gln Glu Ser Ser  His Phe Leu Gln Gly  Ala Lys Lys
    1580                1585                1590

Asn Asn  Leu Ser Leu Ala Ile  Leu Thr Leu Glu Met  Thr Gly Asp
    1595                1600                1605

Gln Arg  Glu Val Gly Ser Leu  Gly Thr Ser Ala Thr  Asn Ser Val
    1610                1615                1620

Thr Tyr  Lys Lys Val Glu Asn  Thr Val Leu Pro Lys  Pro Asp Leu
    1625                1630                1635
```

```
Pro Lys  Thr Ser Gly Lys Val  Glu Leu Leu Pro Lys  Val His Ile
    1640                1645                1650

Tyr Gln  Lys Asp Leu Phe Pro  Thr Glu Thr Ser Asn  Gly Ser Pro
    1655                1660                1665

Gly His  Leu Asp Leu Val Glu  Gly Ser Leu Leu Gln  Gly Thr Glu
    1670                1675                1680

Gly Ala  Ile Lys Trp Asn Glu  Ala Asn Arg Pro Gly  Lys Val Pro
    1685                1690                1695

Phe Leu  Arg Val Ala Thr Glu  Ser Ser Ala Lys Thr  Pro Ser Lys
    1700                1705                1710

Leu Leu  Asp Pro Leu Ala Trp  Asp Asn His Tyr Gly  Thr Gln Ile
    1715                1720                1725

Pro Lys  Glu Glu Trp Lys Ser  Gln Glu Lys Ser Pro  Glu Lys Thr
    1730                1735                1740

Ala Phe  Lys Lys Lys Asp Thr  Ile Leu Ser Leu Asn  Ala Cys Glu
    1745                1750                1755

Ser Asn  His Ala Ile Ala Ala  Ile Asn Glu Gly Gln  Asn Lys Pro
    1760                1765                1770

Glu Ile  Glu Val Thr Trp Ala  Lys Gln Gly Arg Thr  Glu Arg Leu
    1775                1780                1785

Cys Ser  Gln Asn Pro Pro Val  Leu Lys Arg His Gln  Arg Glu Ile
    1790                1795                1800

Thr Arg  Thr Thr Leu Gln Ser  Asp Gln Glu Glu Ile  Asp Tyr Asp
    1805                1810                1815

Asp Thr  Ile Ser Val Glu Met  Lys Lys Glu Asp Phe  Asp Ile Tyr
    1820                1825                1830

Asp Glu  Asp Glu Asn Gln Ser  Pro Arg Ser Phe Gln  Lys Lys Thr
    1835                1840                1845

Arg His  Tyr Phe Ile Ala Ala  Val Glu Arg Leu Trp  Asp Tyr Gly
    1850                1855                1860

Met Ser  Ser Ser Pro His Val  Leu Arg Asn Arg Ala  Gln Ser Gly
    1865                1870                1875

Ser Val  Pro Gln Phe Lys Lys  Val Val Phe Gln Glu  Phe Thr Asp
    1880                1885                1890
```

```
Gly Ser  Phe Thr Gln Pro Leu  Tyr Arg Gly Glu Leu  Asn Glu His
    1895                1900                1905

Leu Gly  Leu Leu Gly Pro Tyr  Ile Arg Ala Glu Val  Glu Asp Asn
    1910                1915                1920

Ile Met  Val Thr Phe Arg Asn  Gln Ala Ser Arg Pro  Tyr Ser Phe
    1925                1930                1935

Tyr Ser  Ser Leu Ile Ser Tyr  Glu Glu Asp Gln Arg  Gln Gly Ala
    1940                1945                1950

Glu Pro  Arg Lys Asn Phe Val  Lys Pro Asn Glu Thr  Lys Thr Tyr
    1955                1960                1965

Phe Trp  Lys Val Gln His His  Met Ala Pro Thr Lys  Asp Glu Phe
    1970                1975                1980

Asp Cys  Lys Ala Trp Ala Tyr  Phe Ser Asp Val Asp  Leu Glu Lys
    1985                1990                1995

Asp Val  His Ser Gly Leu Ile  Gly Pro Leu Leu Val  Cys His Thr
    2000                2005                2010

Asn Thr  Leu Asn Pro Ala His  Gly Arg Gln Val Thr  Val Gln Glu
    2015                2020                2025

Phe Ala  Leu Phe Phe Thr Ile  Phe Asp Glu Thr Lys  Ser Trp Tyr
    2030                2035                2040

Phe Thr  Glu Asn Met Glu Arg  Asn Cys Arg Ala Pro  Cys Asn Ile
    2045                2050                2055

Gln Met  Glu Asp Pro Thr Phe  Lys Glu Asn Tyr Arg  Phe His Ala
    2060                2065                2070

Ile Asn  Gly Tyr Ile Met Asp  Thr Leu Pro Gly Leu  Val Met Ala
    2075                2080                2085

Gln Asp  Gln Arg Ile Arg Trp  Tyr Leu Leu Ser Met  Gly Ser Asn
    2090                2095                2100

Glu Asn  Ile His Ser Ile His  Phe Ser Gly His Val  Phe Thr Val
    2105                2110                2115

Arg Lys  Lys Glu Glu Tyr Lys  Met Ala Leu Tyr Asn  Leu Tyr Pro
    2120                2125                2130

Gly Val  Phe Glu Thr Val Glu  Met Leu Pro Ser Lys  Ala Gly Ile
    2135                2140                2145
```

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Trp Arg  Val Glu Cys Leu Ile  Gly Glu His Leu His  Ala Gly Met
    2150                 2155                 2160

Ser Thr  Leu Phe Leu Val Tyr  Ser Asn Lys Cys Gln  Thr Pro Leu
    2165                 2170                 2175

Gly Met  Ala Ser Gly His Ile  Arg Asp Phe Gln Ile  Thr Ala Ser
    2180                 2185                 2190

Gly Gln  Tyr Gly Gln Trp Ala  Pro Lys Leu Ala Arg  Leu His Tyr
    2195                 2200                 2205

Ser Gly  Ser Ile Asn Ala Trp  Ser Thr Lys Glu Pro  Phe Ser Trp
    2210                 2215                 2220

Ile Lys  Val Asp Leu Leu Ala  Pro Met Ile Ile His  Gly Ile Lys
    2225                 2230                 2235

Thr Gln  Gly Ala Arg Gln Lys  Phe Ser Ser Leu Tyr  Ile Ser Gln
    2240                 2245                 2250

Phe Ile  Ile Met Tyr Ser Leu  Asp Gly Lys Lys Trp  Gln Thr Tyr
    2255                 2260                 2265

Arg Gly  Asn Ser Thr Gly Thr  Leu Met Val Phe Phe  Gly Asn Val
    2270                 2275                 2280

Asp Ser  Ser Gly Ile Lys His  Asn Ile Phe Asn Pro  Pro Ile Ile
    2285                 2290                 2295

Ala Arg  Tyr Ile Arg Leu His  Pro Thr His Tyr Ser  Ile Arg Ser
    2300                 2305                 2310

Thr Leu  Arg Met Glu Leu Met  Gly Cys Asp Leu Asn  Ser Cys Ser
    2315                 2320                 2325

Met Pro  Leu Gly Met Glu Ser  Lys Ala Ile Ser Asp  Ala Gln Ile
    2330                 2335                 2340

Thr Ala  Ser Ser Tyr Phe Thr  Asn Met Phe Ala Thr  Trp Ser Pro
    2345                 2350                 2355

Ser Lys  Ala Arg Leu His Leu  Gln Gly Arg Ser Asn  Ala Trp Arg
    2360                 2365                 2370

Pro Gln  Val Asn Asn Pro Lys  Glu Trp Leu Gln Val  Asp Phe Gln
    2375                 2380                 2385

Lys Thr  Met Lys Val Thr Gly  Val Thr Thr Gln Gly  Val Lys Ser
    2390                 2395                 2400
```

```
Leu Leu  Thr Ser Met Tyr Val  Lys Glu Phe Leu Ile  Ser Ser Ser
    2405                 2410                 2415

Gln Asp  Gly His Gln Trp Thr  Leu Phe Phe Gln Asn  Gly Lys Val
    2420                 2425                 2430

Lys Val  Phe Gln Gly Asn Gln  Asp Ser Phe Thr Pro  Val Val Asn
    2435                 2440                 2445

Ser Leu  Asp Pro Pro Leu Leu  Thr Arg Tyr Leu Arg  Ile His Pro
    2450                 2455                 2460

Gln Ser  Trp Val His Gln Ile  Ala Leu Arg Met Glu  Val Leu Gly
    2465                 2470                 2475

Cys Glu  Ala Gln Asp Leu Tyr  Gly Thr Met Thr Arg  Ile Val Gly
    2480                 2485                 2490

Gly Gly  Ser Thr Ser Glu Ser  Pro Ser Gly Thr Ala  Pro Gly Thr
    2495                 2500                 2505

Ser Pro  Ser Gly Glu Ser Ser  Thr Ala Pro Gly Ser  Thr Ser Glu
    2510                 2515                 2520

Ser Pro  Ser Gly Thr Ala Pro  Gly Ser Thr Ser Glu  Ser Pro Ser
    2525                 2530                 2535

Gly Thr  Ala Pro Gly Thr Ser  Thr Pro Glu Ser Gly  Ser Ala Ser
    2540                 2545                 2550

Pro Gly  Thr Ser Thr Pro Glu  Ser Gly Ser Ala Ser  Pro Gly Ser
    2555                 2560                 2565

Thr Ser  Glu Ser Pro Ser Gly  Thr Ala Pro Gly Ser  Thr Ser Glu
    2570                 2575                 2580

Ser Pro  Ser Gly Thr Ala Pro  Gly Thr Ser Pro Ser  Gly Glu Ser
    2585                 2590                 2595

Ser Thr  Ala Pro Gly Ser Thr  Ser Glu Ser Pro Ser  Gly Thr Ala
    2600                 2605                 2610

Pro Gly  Thr Ser Pro Ser Gly  Glu Ser Ser Thr Ala  Pro Gly Thr
    2615                 2620                 2625

Ser Pro  Ser Gly Glu Ser Ser  Thr Ala Pro Gly Ser  Thr Ser Ser
    2630                 2635                 2640

Thr Ala  Glu Ser Pro Gly Pro  Gly Thr Ser Pro Ser  Gly Glu Ser
    2645                 2650                 2655
```

```
Ser Thr  Ala Pro Gly Thr Ser  Pro Ser Gly Glu Ser  Ser Thr Ala
    2660                 2665                 2670

Pro Gly  Ser Thr Ser Ser Thr  Ala Glu Ser Pro Gly  Pro Gly Thr
    2675                 2680                 2685

Ser Thr  Pro Glu Ser Gly Ser  Ala Ser Pro Gly Thr  Ser Thr Pro
    2690                 2695                 2700

Glu Ser  Gly Ser Ala Ser Pro  Gly Ser Thr Ser Glu  Ser Pro Ser
    2705                 2710                 2715

Gly Thr  Ala Pro Gly Ser Thr  Ser Glu Ser Pro Ser  Gly Thr Ala
    2720                 2725                 2730

Pro Gly  Thr Ser Thr Pro Glu  Ser Gly Ser Ala Ser  Pro Gly Ser
    2735                 2740                 2745

Thr Ser  Ser Thr Ala Glu Ser  Pro Gly Pro Gly Thr  Ser Thr Pro
    2750                 2755                 2760

Glu Ser  Gly Ser Ala Ser Pro  Gly Ser Thr Ser Glu  Ser Pro Ser
    2765                 2770                 2775

Gly Thr  Ala Pro Gly Thr Ser  Pro Ser Gly Glu Ser  Ser Thr Ala
    2780                 2785                 2790

Pro Gly  Ser Thr Ser Ser Thr  Ala Glu Ser Pro Gly  Pro Gly Thr
    2795                 2800                 2805

Ser Pro  Ser Gly Glu Ser Ser  Thr Ala Pro Gly Thr  Ser Thr Pro
    2810                 2815                 2820

Glu Ser  Gly Ser Ala Ser Pro  Gly Ser Thr Ser Ser  Thr Ala Glu
    2825                 2830                 2835

Ser Pro  Gly Pro Gly Ser Thr  Ser Ser Thr Ala Glu  Ser Pro Gly
    2840                 2845                 2850

Pro Gly  Ser Thr Ser Ser Thr  Ala Glu Ser Pro Gly  Pro Gly Ser
    2855                 2860                 2865

Thr Ser  Ser Thr Ala Glu Ser  Pro Gly Pro Gly Thr  Ser Pro Ser
    2870                 2875                 2880

Gly Glu  Ser Ser Thr Ala Pro  Gly Ser Thr Ser Glu  Ser Pro Ser
    2885                 2890                 2895

Gly Thr  Ala Pro Gly Ser Thr  Ser Glu Ser Pro Ser  Gly Thr Ala
    2900                 2905                 2910
```

Pro Gly Thr Ser Thr Pro Glu Ser Gly Pro Xaa Xaa Xaa Gly Ala
2915 2920 2925

Ser Ala Ser Gly Ala Pro Ser Thr Xaa Xaa Xaa Xaa Ser Glu Ser
2930 2935 2940

Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly
2945 2950 2955

Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
2960 2965 2970

Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr
2975 2980 2985

Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Ser Thr Ser Glu Ser
2990 2995 3000

Pro Ser Gly Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser
3005 3010 3015

Ala Ser Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro
3020 3025 3030

Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr
3035 3040 3045

Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro Ser Gly
3050 3055 3060

Glu Ser Ser Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser
3065 3070 3075

Ala Ser Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
3080 3085 3090

Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser
3095 3100 3105

Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser Glu Ser
3110 3115 3120

Pro Ser Gly Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser
3125 3130 3135

Ala Ser Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser Ala Ser Pro
3140 3145 3150

Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro Gly Thr Ser
3155 3160 3165

Thr Pro Glu Ser Gly Ser Ala Ser Pro Gly Ser Thr Ser Ser Thr
3170 3175 3180

Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly
3185 3190 3195

Thr Ala Pro Gly Ser Thr Ser Glu Ser Pro Ser Gly Thr Ala Pro
3200 3205 3210

Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr
3215 3220 3225

Ser Ser Thr Ala Glu Ser Pro Gly Pro Gly Thr Ser Pro Ser Gly
3230 3235 3240

Glu Ser Ser Thr Ala Pro Gly Thr Ser Thr Pro Glu Ser Gly Ser
3245 3250 3255

Ala Ser Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro
3260 3265 3270

Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Thr Ser
3275 3280 3285

Pro Ser Gly Glu Ser Ser Thr Ala Pro Gly Ser Thr Ser Ser Thr
3290 3295 3300

Ala Glu Ser Pro Gly Pro Gly Ser Thr Ser Ser Thr Ala Glu Ser
3305 3310 3315

Pro Gly Pro Gly Thr Ser Pro Ser Gly Glu Ser Ser Thr Ala Pro
3320 3325 3330

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser
3335 3340 3345

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser
3350 3355 3360

Gly Ala Thr Gly Ser Pro
3365

<210> 1580
<211> 3350
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1580
Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
1 5 10 15

```
Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
            20                  25                  30

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
        35                  40                  45

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
    50                  55                  60

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
65                  70                  75                  80

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
                85                  90                  95

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
            100                 105                 110

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
        115                 120                 125

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    130                 135                 140

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
145                 150                 155                 160

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
                165                 170                 175

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            180                 185                 190

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
        195                 200                 205

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
    210                 215                 220

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
225                 230                 235                 240

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
                245                 250                 255

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
            260                 265                 270

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        275                 280                 285
```

```
Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
    290                 295                 300

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
305                 310                 315                 320

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
                325                 330                 335

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
            340                 345                 350

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
        355                 360                 365

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    370                 375                 380

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
385                 390                 395                 400

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
                405                 410                 415

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            420                 425                 430

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
        435                 440                 445

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
    450                 455                 460

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
465                 470                 475                 480

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
                485                 490                 495

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
            500                 505                 510

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
        515                 520                 525

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala
    530                 535                 540

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
545                 550                 555                 560
```

```
Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
                565                 570                 575

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
            580                 585                 590

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
        595                 600                 605

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
    610                 615                 620

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
625                 630                 635                 640

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
                645                 650                 655

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
            660                 665                 670

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
        675                 680                 685

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
    690                 695                 700

Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
705                 710                 715                 720

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
                725                 730                 735

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
            740                 745                 750

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        755                 760                 765

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
    770                 775                 780

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
785                 790                 795                 800

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
                805                 810                 815

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
            820                 825                 830
```

```
Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
        835                 840                 845

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    850                 855                 860

Gly Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp
865                 870                 875                 880

Tyr Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro
                885                 890                 895

Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys
            900                 905                 910

Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys
        915                 920                 925

Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu
    930                 935                 940

Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro
945                 950                 955                 960

Val Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly
                965                 970                 975

Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys
            980                 985                 990

Val Phe Pro Gly Gly Ser His Thr  Tyr Val Trp Gln Val  Leu Lys Glu
        995                 1000                1005

Asn Gly  Pro Met Ala Ser Asp  Pro Leu Cys Leu Thr  Tyr Ser Tyr
    1010                1015                1020

Leu Ser  His Val Asp Leu Val  Lys Asp Leu Asn Ser  Gly Leu Ile
    1025                1030                1035

Gly Ala  Leu Leu Val Cys Arg  Glu Gly Ser Leu Ala  Lys Glu Lys
    1040                1045                1050

Thr Gln  Thr Leu His Lys Phe  Ile Leu Leu Phe Ala  Val Phe Asp
    1055                1060                1065

Glu Gly  Lys Ser Trp His Ser  Glu Thr Lys Asn Ser  Leu Met Gln
    1070                1075                1080

Asp Arg  Asp Ala Ala Ser Ala  Arg Ala Trp Pro Lys  Met His Thr
    1085                1090                1095
```

```
Val Asn  Gly Tyr Val Asn Arg  Ser Leu Pro Gly Leu  Ile Gly Cys
    1100                1105                1110

His Arg  Lys Ser Val Tyr Trp  His Val Ile Gly Met  Gly Thr Thr
    1115                1120                1125

Pro Glu  Val His Ser Ile Phe  Leu Glu Gly His Thr  Phe Leu Val
    1130                1135                1140

Arg Asn  His Arg Gln Ala Ser  Leu Glu Ile Ser Pro  Ile Thr Phe
    1145                1150                1155

Leu Thr  Ala Gln Thr Leu Leu  Met Asp Leu Gly Gln  Phe Leu Leu
    1160                1165                1170

Phe Cys  His Ile Ser Ser His  Gln His Asp Gly Met  Glu Ala Tyr
    1175                1180                1185

Val Lys  Val Asp Ser Cys Pro  Glu Glu Pro Gln Leu  Arg Met Lys
    1190                1195                1200

Asn Asn  Glu Glu Ala Glu Asp  Tyr Asp Asp Asp Leu  Thr Asp Ser
    1205                1210                1215

Glu Met  Asp Val Val Arg Phe  Asp Asp Asp Asn Ser  Pro Ser Phe
    1220                1225                1230

Ile Gln  Ile Arg Ser Val Ala  Lys Lys His Pro Lys  Thr Trp Val
    1235                1240                1245

His Tyr  Ile Ala Ala Glu Glu  Glu Asp Trp Asp Tyr  Ala Pro Leu
    1250                1255                1260

Val Leu  Ala Pro Asp Asp Arg  Ser Tyr Lys Ser Gln  Tyr Leu Asn
    1265                1270                1275

Asn Gly  Pro Gln Arg Ile Gly  Arg Lys Tyr Lys Lys  Val Arg Phe
    1280                1285                1290

Met Ala  Tyr Thr Asp Glu Thr  Phe Lys Thr Arg Glu  Ala Ile Gln
    1295                1300                1305

His Glu  Ser Gly Ile Leu Gly  Pro Leu Leu Tyr Gly  Glu Val Gly
    1310                1315                1320

Asp Thr  Leu Leu Ile Ile Phe  Lys Asn Gln Ala Ser  Arg Pro Tyr
    1325                1330                1335

Asn Ile  Tyr Pro His Gly Ile  Thr Asp Val Arg Pro  Leu Tyr Ser
    1340                1345                1350
```

```
Arg Arg  Leu Pro Lys Gly Val  Lys His Leu Lys Asp  Phe Pro Ile
    1355                1360                1365

Leu Pro  Gly Glu Ile Phe Lys  Tyr Lys Trp Thr Val  Thr Val Glu
    1370                1375                1380

Asp Gly  Pro Thr Lys Ser Asp  Pro Arg Cys Leu Thr  Arg Tyr Tyr
    1385                1390                1395

Ser Ser  Phe Val Asn Met Glu  Arg Asp Leu Ala Ser  Gly Leu Ile
    1400                1405                1410

Gly Pro  Leu Leu Ile Cys Tyr  Lys Glu Ser Val Asp  Gln Arg Gly
    1415                1420                1425

Asn Gln  Ile Met Ser Asp Lys  Arg Asn Val Ile Leu  Phe Ser Val
    1430                1435                1440

Phe Asp  Glu Asn Arg Ser Trp  Tyr Leu Thr Glu Asn  Ile Gln Arg
    1445                1450                1455

Phe Leu  Pro Asn Pro Ala Gly  Val Gln Leu Glu Asp  Pro Glu Phe
    1460                1465                1470

Gln Ala  Ser Asn Ile Met His  Ser Ile Asn Gly Tyr  Val Phe Asp
    1475                1480                1485

Ser Leu  Gln Leu Ser Val Cys  Leu His Glu Val Ala  Tyr Trp Tyr
    1490                1495                1500

Ile Leu  Ser Ile Gly Ala Gln  Thr Asp Phe Leu Ser  Val Phe Phe
    1505                1510                1515

Ser Gly  Tyr Thr Phe Lys His  Lys Met Val Tyr Glu  Asp Thr Leu
    1520                1525                1530

Thr Leu  Phe Pro Phe Ser Gly  Glu Thr Val Phe Met  Ser Met Glu
    1535                1540                1545

Asn Pro  Gly Leu Trp Ile Leu  Gly Cys His Asn Ser  Asp Phe Arg
    1550                1555                1560

Asn Arg  Gly Met Thr Ala Leu  Leu Lys Val Ser Ser  Cys Asp Lys
    1565                1570                1575

Asn Thr  Gly Asp Tyr Tyr Glu  Asp Ser Tyr Glu Asp  Ile Ser Ala
    1580                1585                1590

Tyr Leu  Leu Ser Lys Asn Asn  Ala Ile Glu Pro Arg  Ser Phe Ser
    1595                1600                1605
```

```
Gln Asn  Ser Arg His Pro Ser  Thr Arg Gln Lys Gln  Phe Asn Ala
    1610                 1615                 1620

Thr Thr  Ile Pro Glu Asn Asp  Ile Glu Lys Thr Asp  Pro Trp Phe
    1625                 1630                 1635

Ala His  Arg Thr Pro Met Pro  Lys Ile Gln Asn Val  Ser Ser Ser
    1640                 1645                 1650

Asp Leu  Leu Met Leu Leu Arg  Gln Ser Pro Thr Pro  His Gly Leu
    1655                 1660                 1665

Ser Leu  Ser Asp Leu Gln Glu  Ala Lys Tyr Glu Thr  Phe Ser Asp
    1670                 1675                 1680

Asp Pro  Ser Pro Gly Ala Ile  Asp Ser Asn Asn Ser  Leu Ser Glu
    1685                 1690                 1695

Met Thr  His Phe Arg Pro Gln  Leu His His Ser Gly  Asp Met Val
    1700                 1705                 1710

Phe Thr  Pro Glu Ser Gly Leu  Gln Leu Arg Leu Asn  Glu Lys Leu
    1715                 1720                 1725

Gly Thr  Thr Ala Ala Thr Glu  Leu Lys Lys Leu Asp  Phe Lys Val
    1730                 1735                 1740

Ser Ser  Thr Ser Asn Asn Leu  Ile Ser Thr Ile Pro  Ser Asp Asn
    1745                 1750                 1755

Leu Ala  Ala Gly Thr Asp Asn  Thr Ser Ser Leu Gly  Pro Pro Ser
    1760                 1765                 1770

Met Pro  Val His Tyr Asp Ser  Gln Leu Asp Thr Thr  Leu Phe Gly
    1775                 1780                 1785

Lys Lys  Ser Ser Pro Leu Thr  Glu Ser Gly Gly Pro  Leu Ser Leu
    1790                 1795                 1800

Ser Glu  Glu Asn Asn Asp Ser  Lys Leu Leu Glu Ser  Gly Leu Met
    1805                 1810                 1815

Asn Ser  Gln Glu Ser Ser Trp  Gly Lys Asn Val Ser  Ser Thr Glu
    1820                 1825                 1830

Ser Gly  Arg Leu Phe Lys Gly  Lys Arg Ala His Gly  Pro Ala Leu
    1835                 1840                 1845

Leu Thr  Lys Asp Asn Ala Leu  Phe Lys Val Ser Ile  Ser Leu Leu
    1850                 1855                 1860
```

```
Lys Thr  Asn Lys Thr Ser Asn  Asn Ser Ala Thr Asn  Arg Lys Thr
    1865                1870                1875

His Ile  Asp Gly Pro Ser Leu  Leu Ile Glu Asn Ser  Pro Ser Val
    1880                1885                1890

Trp Gln  Asn Ile Leu Glu Ser  Asp Thr Glu Phe Lys  Lys Val Thr
    1895                1900                1905

Pro Leu  Ile His Asp Arg Met  Leu Met Asp Lys Asn  Ala Thr Ala
    1910                1915                1920

Leu Arg  Leu Asn His Met Ser  Asn Lys Thr Thr Ser  Ser Lys Asn
    1925                1930                1935

Met Glu  Met Val Gln Gln Lys  Lys Glu Gly Pro Ile  Pro Pro Asp
    1940                1945                1950

Ala Gln  Asn Pro Asp Met Ser  Phe Phe Lys Met Leu  Phe Leu Pro
    1955                1960                1965

Glu Ser  Ala Arg Trp Ile Gln  Arg Thr His Gly Lys  Asn Ser Leu
    1970                1975                1980

Asn Ser  Gly Gln Gly Pro Ser  Pro Lys Gln Leu Val  Ser Leu Gly
    1985                1990                1995

Pro Glu  Lys Ser Val Glu Gly  Gln Asn Phe Leu Ser  Glu Lys Asn
    2000                2005                2010

Lys Val  Val Val Gly Lys Gly  Glu Phe Thr Lys Asp  Val Gly Leu
    2015                2020                2025

Lys Glu  Met Val Phe Pro Ser  Ser Arg Asn Leu Phe  Leu Thr Asn
    2030                2035                2040

Leu Asp  Asn Leu His Glu Asn  Asn Thr His Asn Gln  Glu Lys Lys
    2045                2050                2055

Ile Gln  Glu Glu Ile Glu Lys  Lys Glu Thr Leu Ile  Gln Glu Asn
    2060                2065                2070

Val Val  Leu Pro Gln Ile His  Thr Val Thr Gly Thr  Lys Asn Phe
    2075                2080                2085

Met Lys  Asn Leu Phe Leu Leu  Ser Thr Arg Gln Asn  Val Glu Gly
    2090                2095                2100

Ser Tyr  Asp Gly Ala Tyr Ala  Pro Val Leu Gln Asp  Phe Arg Ser
    2105                2110                2115
```

```
Leu Asn  Asp Ser Thr Asn Arg  Thr Lys Lys His Thr  Ala His Phe
    2120                2125                2130

Ser Lys  Lys Gly Glu Glu Glu  Asn Leu Glu Gly Leu  Gly Asn Gln
    2135                2140                2145

Thr Lys  Gln Ile Val Glu Lys  Tyr Ala Cys Thr Thr  Arg Ile Ser
    2150                2155                2160

Pro Asn  Thr Ser Gln Gln Asn  Phe Val Thr Gln Arg  Ser Lys Arg
    2165                2170                2175

Ala Leu  Lys Gln Phe Arg Leu  Pro Leu Glu Glu Thr  Glu Leu Glu
    2180                2185                2190

Lys Arg  Ile Ile Val Asp Asp  Thr Ser Thr Gln Trp  Ser Lys Asn
    2195                2200                2205

Met Lys  His Leu Thr Pro Ser  Thr Leu Thr Gln Ile  Asp Tyr Asn
    2210                2215                2220

Glu Lys  Glu Lys Gly Ala Ile  Thr Gln Ser Pro Leu  Ser Asp Cys
    2225                2230                2235

Leu Thr  Arg Ser His Ser Ile  Pro Gln Ala Asn Arg  Ser Pro Leu
    2240                2245                2250

Pro Ile  Ala Lys Val Ser Ser  Phe Pro Ser Ile Arg  Pro Ile Tyr
    2255                2260                2265

Leu Thr  Arg Val Leu Phe Gln  Asp Asn Ser Ser His  Leu Pro Ala
    2270                2275                2280

Ala Ser  Tyr Arg Lys Lys Asp  Ser Gly Val Gln Glu  Ser Ser His
    2285                2290                2295

Phe Leu  Gln Gly Ala Lys Lys  Asn Asn Leu Ser Leu  Ala Ile Leu
    2300                2305                2310

Thr Leu  Glu Met Thr Gly Asp  Gln Arg Glu Val Gly  Ser Leu Gly
    2315                2320                2325

Thr Ser  Ala Thr Asn Ser Val  Thr Tyr Lys Lys Val  Glu Asn Thr
    2330                2335                2340

Val Leu  Pro Lys Pro Asp Leu  Pro Lys Thr Ser Gly  Lys Val Glu
    2345                2350                2355

Leu Leu  Pro Lys Val His Ile  Tyr Gln Lys Asp Leu  Phe Pro Thr
    2360                2365                2370
```

```
Glu Thr  Ser Asn Gly Ser Pro  Gly His Leu Asp Leu  Val Glu Gly
    2375                 2380                 2385

Ser Leu  Leu Gln Gly Thr Glu  Gly Ala Ile Lys Trp  Asn Glu Ala
    2390                 2395                 2400

Asn Arg  Pro Gly Lys Val Pro  Phe Leu Arg Val Ala  Thr Glu Ser
    2405                 2410                 2415

Ser Ala  Lys Thr Pro Ser Lys  Leu Leu Asp Pro Leu  Ala Trp Asp
    2420                 2425                 2430

Asn His  Tyr Gly Thr Gln Ile  Pro Lys Glu Glu Trp  Lys Ser Gln
    2435                 2440                 2445

Glu Lys  Ser Pro Glu Lys Thr  Ala Phe Lys Lys Lys  Asp Thr Ile
    2450                 2455                 2460

Leu Ser  Leu Asn Ala Cys Glu  Ser Asn His Ala Ile  Ala Ala Ile
    2465                 2470                 2475

Asn Glu  Gly Gln Asn Lys Pro  Glu Ile Glu Val Thr  Trp Ala Lys
    2480                 2485                 2490

Gln Gly  Arg Thr Glu Arg Leu  Cys Ser Gln Asn Pro  Pro Val Leu
    2495                 2500                 2505

Lys Arg  His Gln Arg Glu Ile  Thr Arg Thr Thr Leu  Gln Ser Asp
    2510                 2515                 2520

Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile Ser Val  Glu Met Lys
    2525                 2530                 2535

Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp Glu Asn  Gln Ser Pro
    2540                 2545                 2550

Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr Phe Ile  Ala Ala Val
    2555                 2560                 2565

Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser Ser Pro  His Val Leu
    2570                 2575                 2580

Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro Gln Phe  Lys Lys Val
    2585                 2590                 2595

Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe Thr Gln  Pro Leu Tyr
    2600                 2605                 2610

Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu Leu Gly  Pro Tyr Ile
    2615                 2620                 2625
```

```
Arg Ala  Glu Val Glu Asp Asn  Ile Met Val Thr Phe  Arg Asn Gln
    2630                 2635                 2640

Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser Leu Ile  Ser Tyr Glu
    2645                 2650                 2655

Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg Lys Asn  Phe Val Lys
    2660                 2665                 2670

Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys Val Gln  His His Met
    2675                 2680                 2685

Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys Ala Trp  Ala Tyr Phe
    2690                 2695                 2700

Ser Asp  Val Asp Leu Glu Lys  Asp Val His Ser Gly  Leu Ile Gly
    2705                 2710                 2715

Pro Leu  Leu Val Cys His Thr  Asn Thr Leu Asn Pro  Ala His Gly
    2720                 2725                 2730

Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu Phe Phe  Thr Ile Phe
    2735                 2740                 2745

Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu Asn Met  Glu Arg Asn
    2750                 2755                 2760

Cys Arg  Ala Pro Cys Asn Ile  Gln Met Glu Asp Pro  Thr Phe Lys
    2765                 2770                 2775

Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly Tyr Ile  Met Asp Thr
    2780                 2785                 2790

Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln Arg Ile  Arg Trp Tyr
    2795                 2800                 2805

Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile His Ser  Ile His Phe
    2810                 2815                 2820

Ser Gly  His Val Phe Thr Val  Arg Lys Lys Glu Glu  Tyr Lys Met
    2825                 2830                 2835

Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe Glu Thr  Val Glu Met
    2840                 2845                 2850

Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val Glu Cys  Leu Ile Gly
    2855                 2860                 2865

Glu His  Leu His Ala Gly Met  Ser Thr Leu Phe Leu  Val Tyr Ser
    2870                 2875                 2880
```

```
Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala Ser Gly  His Ile Arg
    2885                 2890                 2895

Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr Gly Gln  Trp Ala Pro
    2900                 2905                 2910

Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser Ile Asn  Ala Trp Ser
    2915                 2920                 2925

Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val Asp Leu  Leu Ala Pro
    2930                 2935                 2940

Met Ile  Ile His Gly Ile Lys  Thr Gln Gly Ala Arg  Gln Lys Phe
    2945                 2950                 2955

Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile Met Tyr  Ser Leu Asp
    2960                 2965                 2970

Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn Ser Thr  Gly Thr Leu
    2975                 2980                 2985

Met Val  Phe Phe Gly Asn Val  Asp Ser Ser Gly Ile  Lys His Asn
    2990                 2995                 3000

Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr Ile Arg  Leu His Pro
    3005                 3010                 3015

Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg Met Glu  Leu Met Gly
    3020                 3025                 3030

Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu Gly Met  Glu Ser Lys
    3035                 3040                 3045

Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser Ser Tyr  Phe Thr Asn
    3050                 3055                 3060

Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala Arg Leu  His Leu Gln
    3065                 3070                 3075

Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val Asn Asn  Pro Lys Glu
    3080                 3085                 3090

Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met Lys Val  Thr Gly Val
    3095                 3100                 3105

Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr Ser Met  Tyr Val Lys
    3110                 3115                 3120

Glu Phe  Leu Ile Ser Ser Ser  Gln Asp Gly His Gln  Trp Thr Leu
    3125                 3130                 3135
```

```
Phe Phe  Gln Asn Gly Lys Val  Lys Val Phe Gln Gly  Asn Gln Asp
    3140                3145                3150

Ser Phe  Thr Pro Val Val Asn  Ser Leu Asp Pro Pro  Leu Leu Thr
    3155                3160                3165

Arg Tyr  Leu Arg Ile His Pro  Gln Ser Trp Val His  Gln Ile Ala
    3170                3175                3180

Leu Arg  Met Glu Val Leu Gly  Cys Glu Ala Gln Asp  Leu Tyr Gly
    3185                3190                3195

Lys Leu  Thr Arg Ala Glu Thr  Gly Gly Ser Glu Pro  Ala Thr Ser
    3200                3205                3210

Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    3215                3220                3225

Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly
    3230                3235                3240

Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr
    3245                3250                3255

Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Ser Glu Pro  Ala Thr Ser
    3260                3265                3270

Gly Ser  Glu Thr Pro Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu
    3275                3280                3285

Thr Pro  Gly Ser Glu Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly
    3290                3295                3300

Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Glu
    3305                3310                3315

Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser
    3320                3325                3330

Gly Ser  Glu Thr Pro Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    3335                3340                3345

Ala Pro
    3350
```

<210> 1581
<211> 2456
<212> PRT
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
polypeptide

<400> 1581
Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
20 25 30

Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
35 40 45

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
50 55 60

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly
65 70 75 80

Ser Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
85 90 95

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
100 105 110

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
115 120 125

Ser Glu Thr Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
130 135 140

Gly Lys Leu Thr Arg Ala Glu Thr Gly Ala Thr Arg Arg Tyr Tyr Leu
145 150 155 160

Gly Ala Val Glu Leu Ser Trp Asp Tyr Met Gln Ser Asp Leu Gly Glu
165 170 175

Leu Pro Val Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe Pro
180 185 190

Phe Asn Thr Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr
195 200 205

Asp His Leu Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly Leu
210 215 220

Leu Gly Pro Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile Thr
225 230 235 240

Leu Lys Asn Met Ala Ser His Pro Val Ser Leu His Ala Val Gly Val
245 250 255

```
Ser Tyr Trp Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser
            260                 265                 270

Gln Arg Glu Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His Thr
        275                 280                 285

Tyr Val Trp Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp Pro
    290                 295                 300

Leu Cys Leu Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys Asp
305                 310                 315                 320

Leu Asn Ser Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly Ser
                325                 330                 335

Leu Ala Lys Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu Phe
            340                 345                 350

Ala Val Phe Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn Ser
        355                 360                 365

Leu Met Gln Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys Met
    370                 375                 380

His Thr Val Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile Gly
385                 390                 395                 400

Cys His Arg Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr Thr
                405                 410                 415

Pro Glu Val His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val Arg
            420                 425                 430

Asn His Arg Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr
        435                 440                 445

Ala Gln Thr Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys His
    450                 455                 460

Ile Ser Ser His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val Asp
465                 470                 475                 480

Ser Cys Pro Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu Ala
                485                 490                 495

Glu Asp Tyr Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val Arg
            500                 505                 510

Phe Asp Asp Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val Ala
        515                 520                 525
```

```
Lys Lys His Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu Glu
    530                 535                 540

Asp Trp Asp Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr
545                 550                 555                 560

Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr
                565                 570                 575

Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg
            580                 585                 590

Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly
        595                 600                 605

Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg
    610                 615                 620

Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu Tyr
625                 630                 635                 640

Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro Ile
                645                 650                 655

Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu Asp
            660                 665                 670

Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser
        675                 680                 685

Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu
    690                 695                 700

Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile Met
705                 710                 715                 720

Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn Arg
                725                 730                 735

Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala
            740                 745                 750

Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met His
        755                 760                 765

Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys Leu
    770                 775                 780

His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp
785                 790                 795                 800
```

```
Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met Val
                805                 810                 815

Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val Phe
            820                 825                 830

Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn Ser
        835                 840                 845

Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser Cys
    850                 855                 860

Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser
865                 870                 875                 880

Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser
                885                 890                 895

Gln Asn Pro Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg Thr
            900                 905                 910

Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser
        915                 920                 925

Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn
    930                 935                 940

Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala
945                 950                 955                 960

Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val
                965                 970                 975

Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val
            980                 985                 990

Val Phe Gln Glu Phe Thr Asp Gly  Ser Phe Thr Gln Pro  Leu Tyr Arg
        995                 1000                1005

Gly Glu  Leu Asn Glu His Leu  Gly Leu Leu Gly Pro  Tyr Ile Arg
    1010                1015                1020

Ala Glu  Val Glu Asp Asn Ile  Met Val Thr Phe Arg  Asn Gln Ala
    1025                1030                1035

Ser Arg  Pro Tyr Ser Phe Tyr  Ser Ser Leu Ile Ser  Tyr Glu Glu
    1040                1045                1050

Asp Gln  Arg Gln Gly Ala Glu  Pro Arg Lys Asn Phe  Val Lys Pro
    1055                1060                1065
```

```
Asn Glu  Thr Lys Thr Tyr Phe  Trp Lys Val Gln His  His Met Ala
    1070                1075                1080

Pro Thr  Lys Asp Glu Phe Asp  Cys Lys Ala Trp Ala  Tyr Phe Ser
    1085                1090                1095

Asp Val  Asp Leu Glu Lys Asp  Val His Ser Gly Leu  Ile Gly Pro
    1100                1105                1110

Leu Leu  Val Cys His Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg
    1115                1120                1125

Gln Val  Thr Val Gln Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp
    1130                1135                1140

Glu Thr  Lys Ser Trp Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys
    1145                1150                1155

Arg Ala  Pro Cys Asn Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu
    1160                1165                1170

Asn Tyr  Arg Phe His Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu
    1175                1180                1185

Pro Gly  Leu Val Met Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu
    1190                1195                1200

Leu Ser  Met Gly Ser Asn Glu  Asn Ile His Ser Ile  His Phe Ser
    1205                1210                1215

Gly His  Val Phe Thr Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala
    1220                1225                1230

Leu Tyr  Asn Leu Tyr Pro Gly  Val Phe Glu Thr Val  Glu Met Leu
    1235                1240                1245

Pro Ser  Lys Ala Gly Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu
    1250                1255                1260

His Leu  His Ala Gly Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn
    1265                1270                1275

Lys Cys  Gln Thr Pro Leu Gly  Met Ala Ser Gly His  Ile Arg Asp
    1280                1285                1290

Phe Gln  Ile Thr Ala Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys
    1295                1300                1305

Leu Ala  Arg Leu His Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr
    1310                1315                1320
```

```
Lys Glu  Pro Phe Ser Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met
    1325                1330                1335

Ile Ile  His Gly Ile Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser
    1340                1345                1350

Ser Leu  Tyr Ile Ser Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly
    1355                1360                1365

Lys Lys  Trp Gln Thr Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met
    1370                1375                1380

Val Phe  Phe Gly Asn Val Asp  Ser Ser Gly Ile Lys  His Asn Ile
    1385                1390                1395

Phe Asn  Pro Pro Ile Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr
    1400                1405                1410

His Tyr  Ser Ile Arg Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys
    1415                1420                1425

Asp Leu  Asn Ser Cys Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala
    1430                1435                1440

Ile Ser  Asp Ala Gln Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met
    1445                1450                1455

Phe Ala  Thr Trp Ser Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly
    1460                1465                1470

Arg Ser  Asn Ala Trp Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp
    1475                1480                1485

Leu Gln  Val Asp Phe Gln Lys  Thr Met Lys Val Thr  Gly Val Thr
    1490                1495                1500

Thr Gln  Gly Val Lys Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu
    1505                1510                1515

Phe Leu  Ile Ser Ser Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe
    1520                1525                1530

Phe Gln  Asn Gly Lys Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser
    1535                1540                1545

Phe Thr  Pro Val Val Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg
    1550                1555                1560

Tyr Leu  Arg Ile His Pro Gln  Ser Trp Val His Gln  Ile Ala Leu
    1565                1570                1575
```

```
Arg Met  Glu Val Leu Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Gly
    1580                 1585                 1590

Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Glu
    1595                 1600                 1605

Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser
    1610                 1615                 1620

Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr
    1625                 1630                 1635

Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    1640                 1645                 1650

Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Glu
    1655                 1660                 1665

Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser
    1670                 1675                 1680

Gly Ser  Glu Thr Pro Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu
    1685                 1690                 1695

Thr Pro  Gly Ser Pro Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly
    1700                 1705                 1710

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Thr
    1715                 1720                 1725

Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser
    1730                 1735                 1740

Glu Gly  Ser Ala Pro Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr
    1745                 1750                 1755

Glu Glu  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    1760                 1765                 1770

Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Glu
    1775                 1780                 1785

Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser
    1790                 1795                 1800

Glu Gly  Ser Ala Pro Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1805                 1810                 1815

Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly
    1820                 1825                 1830
```

```
Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr
    1835                1840                1845

Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr Ser Glu  Ser Ala Thr
    1850                1855                1860

Pro Glu  Ser Gly Pro Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    1865                1870                1875

Gly Pro  Gly Ser Pro Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly
    1880                1885                1890

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro
    1895                1900                1905

Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr
    1910                1915                1920

Pro Glu  Ser Gly Pro Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    1925                1930                1935

Ala Pro  Gly Thr Ser Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly
    1940                1945                1950

Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr
    1955                1960                1965

Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser
    1970                1975                1980

Glu Gly  Ser Ala Pro Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    1985                1990                1995

Ala Pro  Gly Ser Pro Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly
    2000                2005                2010

Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Glu
    2015                2020                2025

Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser
    2030                2035                2040

Gly Ser  Glu Thr Pro Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    2045                2050                2055

Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly
    2060                2065                2070

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Thr
    2075                2080                2085
```

```
Glu Pro  Ser Glu Gly Ser Ala  Pro Gly Thr Ser Glu  Ser Ala Thr
    2090                 2095                 2100

Pro Glu  Ser Gly Pro Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr
    2105                 2110                 2115

Glu Glu  Gly Ser Pro Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly
    2120                 2125                 2130

Ser Pro  Ala Gly Ser Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Glu
    2135                 2140                 2145

Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser
    2150                 2155                 2160

Glu Gly  Ser Ala Pro Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser
    2165                 2170                 2175

Gly Pro  Gly Ser Glu Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly
    2180                 2185                 2190

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro
    2195                 2200                 2205

Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr
    2210                 2215                 2220

Pro Glu  Ser Gly Pro Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser
    2225                 2230                 2235

Ala Pro  Gly Ser Pro Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly
    2240                 2245                 2250

Thr Ser  Glu Ser Ala Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro
    2255                 2260                 2265

Ala Thr  Ser Gly Ser Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr
    2270                 2275                 2280

Pro Glu  Ser Gly Pro Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr
    2285                 2290                 2295

Glu Glu  Gly Ser Pro Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly
    2300                 2305                 2310

Thr Ser  Thr Glu Pro Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Glu
    2315                 2320                 2325

Ser Ala  Thr Pro Glu Ser Gly  Pro Gly Thr Ser Glu  Ser Ala Thr
    2330                 2335                 2340
```

Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser
2345 2350 2355

Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
2360 2365 2370

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
2375 2380 2385

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser
2390 2395 2400

Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser
2405 2410 2415

Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
2420 2425 2430

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
2435 2440 2445

Glu Pro Ser Glu Gly Ser Ala Pro
2450 2455

<210> 1582
<211> 2361
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1582
Met Ala Glu Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Pro
1 5 10 15

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
20 25 30

Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
35 40 45

Pro Gly Lys Leu Thr Arg Ala Glu Thr Gly Ala Thr Arg Arg Tyr Tyr
50 55 60

Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr Met Gln Ser Asp Leu Gly
65 70 75 80

Glu Leu Pro Val Asp Ala Arg Phe Pro Pro Arg Val Pro Lys Ser Phe
85 90 95

Pro Phe Asn Thr Ser Val Val Tyr Lys Lys Thr Leu Phe Val Glu Phe
100 105 110

```
Thr Asp His Leu Phe Asn Ile Ala Lys Pro Arg Pro Pro Trp Met Gly
        115                 120                 125

Leu Leu Gly Pro Thr Ile Gln Ala Glu Val Tyr Asp Thr Val Val Ile
    130                 135                 140

Thr Leu Lys Asn Met Ala Ser His Pro Val Ser Leu His Ala Val Gly
145                 150                 155                 160

Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala Glu Tyr Asp Asp Gln Thr
                165                 170                 175

Ser Gln Arg Glu Lys Glu Asp Asp Lys Val Phe Pro Gly Gly Ser His
            180                 185                 190

Thr Tyr Val Trp Gln Val Leu Lys Glu Asn Gly Pro Met Ala Ser Asp
        195                 200                 205

Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser His Val Asp Leu Val Lys
    210                 215                 220

Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu Leu Val Cys Arg Glu Gly
225                 230                 235                 240

Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu His Lys Phe Ile Leu Leu
                245                 250                 255

Phe Ala Val Phe Asp Glu Gly Lys Ser Trp His Ser Glu Thr Lys Asn
            260                 265                 270

Ser Leu Met Gln Asp Arg Asp Ala Ala Ser Ala Arg Ala Trp Pro Lys
        275                 280                 285

Met His Thr Val Asn Gly Tyr Val Asn Arg Ser Leu Pro Gly Leu Ile
    290                 295                 300

Gly Cys His Arg Lys Ser Val Tyr Trp His Val Ile Gly Met Gly Thr
305                 310                 315                 320

Thr Pro Glu Val His Ser Ile Phe Leu Glu Gly His Thr Phe Leu Val
                325                 330                 335

Arg Asn His Arg Gln Ala Ser Leu Glu Ile Ser Pro Ile Thr Phe Leu
            340                 345                 350

Thr Ala Gln Thr Leu Leu Met Asp Leu Gly Gln Phe Leu Leu Phe Cys
        355                 360                 365

His Ile Ser Ser His Gln His Asp Gly Met Glu Ala Tyr Val Lys Val
    370                 375                 380
```

```
Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg Met Lys Asn Asn Glu Glu
385                 390                 395                 400

Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp Ser Glu Met Asp Val Val
                405                 410                 415

Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe Ile Gln Ile Arg Ser Val
            420                 425                 430

Ala Lys Lys His Pro Lys Thr Trp Val His Tyr Ile Ala Ala Glu Glu
        435                 440                 445

Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu Ala Pro Asp Asp Arg Ser
    450                 455                 460

Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro Gln Arg Ile Gly Arg Lys
465                 470                 475                 480

Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr Asp Glu Thr Phe Lys Thr
                485                 490                 495

Arg Glu Ala Ile Gln His Glu Ser Gly Ile Leu Gly Pro Leu Leu Tyr
            500                 505                 510

Gly Glu Val Gly Asp Thr Leu Leu Ile Ile Phe Lys Asn Gln Ala Ser
        515                 520                 525

Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile Thr Asp Val Arg Pro Leu
    530                 535                 540

Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys His Leu Lys Asp Phe Pro
545                 550                 555                 560

Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys Trp Thr Val Thr Val Glu
                565                 570                 575

Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys Leu Thr Arg Tyr Tyr Ser
            580                 585                 590

Ser Phe Val Asn Met Glu Arg Asp Leu Ala Ser Gly Leu Ile Gly Pro
        595                 600                 605

Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp Gln Arg Gly Asn Gln Ile
    610                 615                 620

Met Ser Asp Lys Arg Asn Val Ile Leu Phe Ser Val Phe Asp Glu Asn
625                 630                 635                 640

Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln Arg Phe Leu Pro Asn Pro
                645                 650                 655
```

```
Ala Gly Val Gln Leu Glu Asp Pro Glu Phe Gln Ala Ser Asn Ile Met
            660                 665                 670

His Ser Ile Asn Gly Tyr Val Phe Asp Ser Leu Gln Leu Ser Val Cys
        675                 680                 685

Leu His Glu Val Ala Tyr Trp Tyr Ile Leu Ser Ile Gly Ala Gln Thr
    690                 695                 700

Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr Thr Phe Lys His Lys Met
705                 710                 715                 720

Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro Phe Ser Gly Glu Thr Val
                725                 730                 735

Phe Met Ser Met Glu Asn Pro Gly Leu Trp Ile Leu Gly Cys His Asn
            740                 745                 750

Ser Asp Phe Arg Asn Arg Gly Met Thr Ala Leu Leu Lys Val Ser Ser
        755                 760                 765

Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile
    770                 775                 780

Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala Ile Glu Pro Arg Ser Phe
785                 790                 795                 800

Ser Gln Asn Pro Pro Val Leu Lys Arg His Gln Arg Glu Ile Thr Arg
                805                 810                 815

Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile
            820                 825                 830

Ser Val Glu Met Lys Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu
        835                 840                 845

Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile
    850                 855                 860

Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His
865                 870                 875                 880

Val Leu Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys
                885                 890                 895

Val Val Phe Gln Glu Phe Thr Asp Gly Ser Phe Thr Gln Pro Leu Tyr
            900                 905                 910

Arg Gly Glu Leu Asn Glu His Leu Gly Leu Leu Gly Pro Tyr Ile Arg
        915                 920                 925
```

```
Ala Glu Val Glu Asp Asn Ile Met Val Thr Phe Arg Asn Gln Ala Ser
    930                 935                 940

Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile Ser Tyr Glu Glu Asp Gln
945                 950                 955                 960

Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe Val Lys Pro Asn Glu Thr
                965                 970                 975

Lys Thr Tyr Phe Trp Lys Val Gln His His Met Ala Pro Thr Lys Asp
            980                 985                 990

Glu Phe Asp Cys Lys Ala Trp Ala  Tyr Phe Ser Asp Val  Asp Leu Glu
        995                 1000                1005

Lys Asp  Val His Ser Gly Leu  Ile Gly Pro Leu Leu  Val Cys His
    1010                1015                1020

Thr Asn  Thr Leu Asn Pro Ala  His Gly Arg Gln Val  Thr Val Gln
    1025                1030                1035

Glu Phe  Ala Leu Phe Phe Thr  Ile Phe Asp Glu Thr  Lys Ser Trp
    1040                1045                1050

Tyr Phe  Thr Glu Asn Met Glu  Arg Asn Cys Arg Ala  Pro Cys Asn
    1055                1060                1065

Ile Gln  Met Glu Asp Pro Thr  Phe Lys Glu Asn Tyr  Arg Phe His
    1070                1075                1080

Ala Ile  Asn Gly Tyr Ile Met  Asp Thr Leu Pro Gly  Leu Val Met
    1085                1090                1095

Ala Gln  Asp Gln Arg Ile Arg  Trp Tyr Leu Leu Ser  Met Gly Ser
    1100                1105                1110

Asn Glu  Asn Ile His Ser Ile  His Phe Ser Gly His  Val Phe Thr
    1115                1120                1125

Val Arg  Lys Lys Glu Glu Tyr  Lys Met Ala Leu Tyr  Asn Leu Tyr
    1130                1135                1140

Pro Gly  Val Phe Glu Thr Val  Glu Met Leu Pro Ser  Lys Ala Gly
    1145                1150                1155

Ile Trp  Arg Val Glu Cys Leu  Ile Gly Glu His Leu  His Ala Gly
    1160                1165                1170

Met Ser  Thr Leu Phe Leu Val  Tyr Ser Asn Lys Cys  Gln Thr Pro
    1175                1180                1185
```

```
Leu Gly  Met Ala Ser Gly His  Ile Arg Asp Phe Gln  Ile Thr Ala
    1190                 1195                 1200

Ser Gly  Gln Tyr Gly Gln Trp  Ala Pro Lys Leu Ala  Arg Leu His
    1205                 1210                 1215

Tyr Ser  Gly Ser Ile Asn Ala  Trp Ser Thr Lys Glu  Pro Phe Ser
    1220                 1225                 1230

Trp Ile  Lys Val Asp Leu Leu  Ala Pro Met Ile Ile  His Gly Ile
    1235                 1240                 1245

Lys Thr  Gln Gly Ala Arg Gln  Lys Phe Ser Ser Leu  Tyr Ile Ser
    1250                 1255                 1260

Gln Phe  Ile Ile Met Tyr Ser  Leu Asp Gly Lys Lys  Trp Gln Thr
    1265                 1270                 1275

Tyr Arg  Gly Asn Ser Thr Gly  Thr Leu Met Val Phe  Phe Gly Asn
    1280                 1285                 1290

Val Asp  Ser Ser Gly Ile Lys  His Asn Ile Phe Asn  Pro Pro Ile
    1295                 1300                 1305

Ile Ala  Arg Tyr Ile Arg Leu  His Pro Thr His Tyr  Ser Ile Arg
    1310                 1315                 1320

Ser Thr  Leu Arg Met Glu Leu  Met Gly Cys Asp Leu  Asn Ser Cys
    1325                 1330                 1335

Ser Met  Pro Leu Gly Met Glu  Ser Lys Ala Ile Ser  Asp Ala Gln
    1340                 1345                 1350

Ile Thr  Ala Ser Ser Tyr Phe  Thr Asn Met Phe Ala  Thr Trp Ser
    1355                 1360                 1365

Pro Ser  Lys Ala Arg Leu His  Leu Gln Gly Arg Ser  Asn Ala Trp
    1370                 1375                 1380

Arg Pro  Gln Val Asn Asn Pro  Lys Glu Trp Leu Gln  Val Asp Phe
    1385                 1390                 1395

Gln Lys  Thr Met Lys Val Thr  Gly Val Thr Thr Gln  Gly Val Lys
    1400                 1405                 1410

Ser Leu  Leu Thr Ser Met Tyr  Val Lys Glu Phe Leu  Ile Ser Ser
    1415                 1420                 1425

Ser Gln  Asp Gly His Gln Trp  Thr Leu Phe Phe Gln  Asn Gly Lys
    1430                 1435                 1440
```

```
Val Lys  Val Phe Gln Gly Asn  Gln Asp Ser Phe Thr  Pro Val Val
    1445                1450                1455

Asn Ser  Leu Asp Pro Pro Leu  Leu Thr Arg Tyr Leu  Arg Ile His
    1460                1465                1470

Pro Gln  Ser Trp Val His Gln  Ile Ala Leu Arg Met  Glu Val Leu
    1475                1480                1485

Gly Cys  Glu Ala Gln Asp Leu  Tyr Gly Gly Ser Pro  Ala Gly Ser
    1490                1495                1500

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1505                1510                1515

Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1520                1525                1530

Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser
    1535                1540                1545

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Thr Glu Pro
    1550                1555                1560

Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1565                1570                1575

Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro
    1580                1585                1590

Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Ser Pro
    1595                1600                1605

Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Thr Ser  Glu Ser Ala
    1610                1615                1620

Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1625                1630                1635

Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1640                1645                1650

Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr Glu Glu  Gly Thr Ser
    1655                1660                1665

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Thr Glu Pro
    1670                1675                1680

Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1685                1690                1695
```

```
Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1700                1705                1710

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    1715                1720                1725

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Thr Glu Pro
    1730                1735                1740

Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1745                1750                1755

Ser Ala  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1760                1765                1770

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Pro
    1775                1780                1785

Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Thr Ser  Glu Ser Ala
    1790                1795                1800

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1805                1810                1815

Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1820                1825                1830

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1835                1840                1845

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Thr Glu Pro
    1850                1855                1860

Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1865                1870                1875

Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1880                1885                1890

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Ser Pro
    1895                1900                1905

Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Thr Ser  Thr Glu Pro
    1910                1915                1920

Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    1925                1930                1935

Ser Gly  Pro Gly Ser Glu Pro  Ala Thr Ser Gly Ser  Glu Thr Pro
    1940                1945                1950
```

```
Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    1955                 1960                 1965

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1970                 1975                 1980

Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1985                 1990                 1995

Ser Ala  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    2000                 2005                 2010

Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr Glu Glu  Gly Ser Pro
    2015                 2020                 2025

Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Ser Pro  Ala Gly Ser
    2030                 2035                 2040

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    2045                 2050                 2055

Ser Gly  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    2060                 2065                 2070

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    2075                 2080                 2085

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    2090                 2095                 2100

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    2105                 2110                 2115

Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    2120                 2125                 2130

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Ser Pro
    2135                 2140                 2145

Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Thr Ser  Glu Ser Ala
    2150                 2155                 2160

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    2165                 2170                 2175

Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    2180                 2185                 2190

Gly Ser  Pro Ala Gly Ser Pro  Thr Ser Thr Glu Glu  Gly Ser Pro
    2195                 2200                 2205
```

```
Ala Gly  Ser Pro Thr Ser Thr  Glu Glu Gly Thr Ser  Thr Glu Pro
    2210                 2215                 2220


Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Glu Ser Ala  Thr Pro Glu
    2225                 2230                 2235


Ser Gly  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    2240                 2245                 2250


Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    2255                 2260                 2265


Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Ser Glu  Pro Ala Thr
    2270                 2275                 2280


Ser Gly  Ser Glu Thr Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    2285                 2290                 2295


Thr Glu  Glu Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    2300                 2305                 2310


Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Ser Glu
    2315                 2320                 2325


Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    2330                 2335                 2340


Thr Pro  Glu Ser Gly Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    2345                 2350                 2355


Ser Ala  Pro
    2360


<210> 1583
<211> 1734
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
      polypeptide

<400> 1583
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1               5                  10                  15


Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30


Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45
```

```
Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320
```

```
Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590
```

```
Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860
```

```
Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1025                1030                1035

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1040                1045                1050

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1055                1060                1065

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1070                1075                1080

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1085                1090                1095

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1100                1105                1110

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1115                1120                1125
```

```
Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1130                1135                1140

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1145                1150                1155

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1160                1165                1170

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1175                1180                1185

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1190                1195                1200

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1205                1210                1215

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1220                1225                1230

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1235                1240                1245

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1250                1255                1260

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1265                1270                1275

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1280                1285                1290

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1295                1300                1305

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1310                1315                1320

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1325                1330                1335

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1340                1345                1350

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1355                1360                1365

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1370                1375                1380
```

```
Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1385                1390                1395

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1400                1405                1410

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1415                1420                1425

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Lys Leu  Thr Arg Ala
    1430                1435                1440

Glu Thr  Gly Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    1445                1450                1455

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Thr
    1460                1465                1470

Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro
    1475                1480                1485

Ser Gly  Ala Thr Gly Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala
    1490                1495                1500

Ser Ser  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1505                1510                1515

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser
    1520                1525                1530

Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Ser  Ser Thr Pro
    1535                1540                1545

Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1550                1555                1560

Gly Thr  Gly Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
    1565                1570                1575

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Thr
    1580                1585                1590

Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro
    1595                1600                1605

Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1610                1615                1620

Gly Thr  Gly Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
    1625                1630                1635
```

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
1640 1645 1650

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro
1655 1660 1665

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
1670 1675 1680

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
1685 1690 1695

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr
1700 1705 1710

Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
1715 1720 1725

Ser Gly Ala Thr Gly Ser
1730

<210> 1584
<211> 2310
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1584
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

```
Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400
```

```
Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670
```

```
Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940
```

```
Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1025                1030                1035

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1040                1045                1050

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1055                1060                1065

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1070                1075                1080

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1085                1090                1095

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1100                1105                1110

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1115                1120                1125

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1130                1135                1140

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1145                1150                1155

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1160                1165                1170

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1175                1180                1185

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1190                1195                1200
```

```
Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1205                 1210                 1215

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1220                 1225                 1230

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1235                 1240                 1245

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1250                 1255                 1260

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1265                 1270                 1275

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1280                 1285                 1290

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1295                 1300                 1305

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1310                 1315                 1320

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1325                 1330                 1335

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1340                 1345                 1350

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1355                 1360                 1365

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1370                 1375                 1380

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1385                 1390                 1395

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1400                 1405                 1410

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1415                 1420                 1425

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Lys Leu  Thr Arg Ala
    1430                 1435                 1440

Glu Thr  Gly Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1445                 1450                 1455
```

```
Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr Gly Pro  Gly Ser Ser
    1460                1465                1470

Pro Ser  Ala Ser Thr Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly
    1475                1480                1485

Thr Ala  Ser Ser Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    1490                1495                1500

Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro
    1505                1510                1515

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Thr Pro
    1520                1525                1530

Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly Ser Ser  Thr Pro Ser
    1535                1540                1545

Gly Ala  Thr Gly Ser Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    1550                1555                1560

Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1565                1570                1575

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Thr Pro
    1580                1585                1590

Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly Ser Ser  Thr Pro Ser
    1595                1600                1605

Gly Ala  Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
    1610                1615                1620

Gly Ser  Pro Gly Thr Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro
    1625                1630                1635

Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser
    1640                1645                1650

Pro Ser  Ala Ser Thr Gly Thr  Gly Pro Gly Ser Ser  Pro Ser Ala
    1655                1660                1665

Ser Thr  Gly Thr Gly Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    1670                1675                1680

Gly Ser  Pro Gly Ser Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro
    1685                1690                1695

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1700                1705                1710
```

```
Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    1715                1720                1725

Ser Ser  Thr Gly Ser Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    1730                1735                1740

Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1745                1750                1755

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1760                1765                1770

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala
    1775                1780                1785

Ser Thr  Gly Thr Gly Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    1790                1795                1800

Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1805                1810                1815

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1820                1825                1830

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser
    1835                1840                1845

Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    1850                1855                1860

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1865                1870                1875

Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser Ser Pro  Gly Ser Ser
    1880                1885                1890

Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser
    1895                1900                1905

Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    1910                1915                1920

Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro
    1925                1930                1935

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1940                1945                1950

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Thr Pro  Gly Ser Gly
    1955                1960                1965
```

```
Thr Ala  Ser Ser Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
    1970                 1975                 1980

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1985                 1990                 1995

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    2000                 2005                 2010

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Thr Pro  Gly Ser Gly
    2015                 2020                 2025

Thr Ala  Ser Ser Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    2030                 2035                 2040

Gly Ser  Pro Gly Thr Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro
    2045                 2050                 2055

Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Gly Thr Pro
    2060                 2065                 2070

Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly Ser Ser  Thr Pro Ser
    2075                 2080                 2085

Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    2090                 2095                 2100

Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro
    2105                 2110                 2115

Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr Gly Pro  Gly Ala Ser
    2120                 2125                 2130

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Thr Pro  Gly Ser Gly
    2135                 2140                 2145

Thr Ala  Ser Ser Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    2150                 2155                 2160

Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro
    2165                 2170                 2175

Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr Gly Pro  Gly Ala Ser
    2180                 2185                 2190

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    2195                 2200                 2205

Ser Ser  Thr Gly Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    2210                 2215                 2220
```

```
Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro
    2225                 2230                 2235


Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ser Ser
    2240                 2245                 2250


Pro Ser  Ala Ser Thr Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly
    2255                 2260                 2265


Thr Ala  Ser Ser Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    2270                 2275                 2280


Gly Ser  Pro Gly Ser Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro
    2285                 2290                 2295


Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro
    2300                 2305                 2310


<210> 1585
<211> 2599
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
      polypeptide

<400> 1585
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1               5                  10                  15


Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30


Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45


Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60


Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80


Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95


Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110


Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125
```

```
Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400
```

```
Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670
```

```
Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Pro Gly Ala Ser Pro Gly
            740                 745                 750

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
        755                 760                 765

Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
    770                 775                 780

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser
785                 790                 795                 800

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
                805                 810                 815

Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
            820                 825                 830

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
        835                 840                 845

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
    850                 855                 860

Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
865                 870                 875                 880

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser
                885                 890                 895

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
            900                 905                 910

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
        915                 920                 925

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly
    930                 935                 940
```

```
Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
945                 950                 955                 960

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
                965                 970                 975

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly
            980                 985                 990

Thr Ser Ser Thr Gly Ser Pro Gly  Ser Ser Pro Ser Ala  Ser Thr Gly
        995                 1000                1005

Thr Gly  Pro Gly Thr Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro
    1010                1015                1020

Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Pro Val Leu
    1025                1030                1035

Lys Arg  His Gln Arg Glu Ile  Thr Arg Thr Thr Leu  Gln Ser Asp
    1040                1045                1050

Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile Ser Val  Glu Met Lys
    1055                1060                1065

Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp Glu Asn  Gln Ser Pro
    1070                1075                1080

Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr Phe Ile  Ala Ala Val
    1085                1090                1095

Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser Ser Pro  His Val Leu
    1100                1105                1110

Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro Gln Phe  Lys Lys Val
    1115                1120                1125

Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe Thr Gln  Pro Leu Tyr
    1130                1135                1140

Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu Leu Gly  Pro Tyr Ile
    1145                1150                1155

Arg Ala  Glu Val Glu Asp Asn  Ile Met Val Thr Phe  Arg Asn Gln
    1160                1165                1170

Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser Leu Ile  Ser Tyr Glu
    1175                1180                1185

Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg Lys Asn  Phe Val Lys
    1190                1195                1200
```

```
Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys Val Gln  His His Met
    1205                1210                1215

Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys Ala Trp  Ala Tyr Phe
    1220                1225                1230

Ser Asp  Val Asp Leu Glu Lys  Asp Val His Ser Gly  Leu Ile Gly
    1235                1240                1245

Pro Leu  Leu Val Cys His Thr  Asn Thr Leu Asn Pro  Ala His Gly
    1250                1255                1260

Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu Phe Phe  Thr Ile Phe
    1265                1270                1275

Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu Asn Met  Glu Arg Asn
    1280                1285                1290

Cys Arg  Ala Pro Cys Asn Ile  Gln Met Glu Asp Pro  Thr Phe Lys
    1295                1300                1305

Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly Tyr Ile  Met Asp Thr
    1310                1315                1320

Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln Arg Ile  Arg Trp Tyr
    1325                1330                1335

Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile His Ser  Ile His Phe
    1340                1345                1350

Ser Gly  His Val Phe Thr Val  Arg Lys Lys Glu Glu  Tyr Lys Met
    1355                1360                1365

Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe Glu Thr  Val Glu Met
    1370                1375                1380

Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val Glu Cys  Leu Ile Gly
    1385                1390                1395

Glu His  Leu His Ala Gly Met  Ser Thr Leu Phe Leu  Val Tyr Ser
    1400                1405                1410

Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala Ser Gly  His Ile Arg
    1415                1420                1425

Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr Gly Gln  Trp Ala Pro
    1430                1435                1440

Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser Ile Asn  Ala Trp Ser
    1445                1450                1455
```

```
Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val Asp Leu  Leu Ala Pro
    1460                1465                1470

Met Ile  Ile His Gly Ile Lys  Thr Gln Gly Ala Arg  Gln Lys Phe
    1475                1480                1485

Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile Met Tyr  Ser Leu Asp
    1490                1495                1500

Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn Ser Thr  Gly Thr Leu
    1505                1510                1515

Met Val  Phe Phe Gly Asn Val  Asp Ser Ser Gly Ile  Lys His Asn
    1520                1525                1530

Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr Ile Arg  Leu His Pro
    1535                1540                1545

Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg Met Glu  Leu Met Gly
    1550                1555                1560

Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu Gly Met  Glu Ser Lys
    1565                1570                1575

Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser Ser Tyr  Phe Thr Asn
    1580                1585                1590

Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala Arg Leu  His Leu Gln
    1595                1600                1605

Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val Asn Asn  Pro Lys Glu
    1610                1615                1620

Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met Lys Val  Thr Gly Val
    1625                1630                1635

Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr Ser Met  Tyr Val Lys
    1640                1645                1650

Glu Phe  Leu Ile Ser Ser Ser  Gln Asp Gly His Gln  Trp Thr Leu
    1655                1660                1665

Phe Phe  Gln Asn Gly Lys Val  Lys Val Phe Gln Gly  Asn Gln Asp
    1670                1675                1680

Ser Phe  Thr Pro Val Val Asn  Ser Leu Asp Pro Pro  Leu Leu Thr
    1685                1690                1695

Arg Tyr  Leu Arg Ile His Pro  Gln Ser Trp Val His  Gln Ile Ala
    1700                1705                1710
```

```
Leu Arg  Met Glu Val Leu Gly  Cys Glu Ala Gln Asp  Leu Tyr Lys
    1715                1720                1725

Leu Thr  Arg Ala Glu Thr Gly  Gly Ala Ser Pro Gly  Thr Ser Ser
    1730                1735                1740

Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
    1745                1750                1755

Pro Gly  Ser Ser Pro Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Thr
    1760                1765                1770

Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro
    1775                1780                1785

Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1790                1795                1800

Gly Thr  Gly Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    1805                1810                1815

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser
    1820                1825                1830

Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Thr  Pro Gly Ser
    1835                1840                1845

Gly Thr  Ala Ser Ser Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1850                1855                1860

Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    1865                1870                1875

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser
    1880                1885                1890

Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1895                1900                1905

Thr Ser  Ser Thr Gly Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala
    1910                1915                1920

Ser Ser  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1925                1930                1935

Pro Gly  Ser Ser Pro Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ser
    1940                1945                1950

Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro Gly Ser  Ser Thr Pro
    1955                1960                1965
```

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
1970 1975 1980

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
1985 1990 1995

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
2000 2005 2010

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser
2015 2020 2025

Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
2030 2035 2040

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
2045 2050 2055

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser
2060 2065 2070

Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser
2075 2080 2085

Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
2090 2095 2100

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
2105 2110 2115

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser
2120 2125 2130

Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
2135 2140 2145

Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
2150 2155 2160

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
2165 2170 2175

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser
2180 2185 2190

Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
2195 2200 2205

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
2210 2215 2220

```
Gly Thr  Gly Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    2225                 2230                 2235

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Thr
    2240                 2245                 2250

Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro Gly Ala  Ser Pro Gly
    2255                 2260                 2265

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    2270                 2275                 2280

Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    2285                 2290                 2295

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Thr
    2300                 2305                 2310

Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro
    2315                 2320                 2325

Ser Gly  Ala Thr Gly Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala
    2330                 2335                 2340

Ser Ser  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    2345                 2350                 2355

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser
    2360                 2365                 2370

Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Ser  Ser Thr Pro
    2375                 2380                 2385

Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    2390                 2395                 2400

Gly Thr  Gly Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
    2405                 2410                 2415

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Thr
    2420                 2425                 2430

Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro
    2435                 2440                 2445

Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    2450                 2455                 2460

Gly Thr  Gly Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
    2465                 2470                 2475
```

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
2480 2485 2490

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro
2495 2500 2505

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
2510 2515 2520

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
2525 2530 2535

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr
2540 2545 2550

Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
2555 2560 2565

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
2570 2575 2580

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
2585 2590 2595

Pro

<210> 1586
<211> 2599
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1586
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

```
Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365
```

```
Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640
```

```
Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gly Pro Gly Ala Ser Pro Gly Thr Ser
            740                 745                 750

Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
        755                 760                 765

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser
    770                 775                 780

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
785                 790                 795                 800

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
                805                 810                 815

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser
            820                 825                 830

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly
        835                 840                 845

Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
    850                 855                 860

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
865                 870                 875                 880

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
                885                 890                 895

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
            900                 905                 910
```

```
Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser
        915                 920                 925

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser
    930                 935                 940

Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
945                 950                 955                 960

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
                965                 970                 975

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser
            980                 985                 990

Ser Thr Gly Ser Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
        995                 1000                1005

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser
    1010                1015                1020

Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Gln Asn Pro  Pro Val Leu
    1025                1030                1035

Lys Arg  His Gln Arg Glu Ile  Thr Arg Thr Thr Leu  Gln Ser Asp
    1040                1045                1050

Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile Ser Val  Glu Met Lys
    1055                1060                1065

Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp Glu Asn  Gln Ser Pro
    1070                1075                1080

Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr Phe Ile  Ala Ala Val
    1085                1090                1095

Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser Ser Pro  His Val Leu
    1100                1105                1110

Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro Gln Phe  Lys Lys Val
    1115                1120                1125

Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe Thr Gln  Pro Leu Tyr
    1130                1135                1140

Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu Leu Gly  Pro Tyr Ile
    1145                1150                1155

Arg Ala  Glu Val Glu Asp Asn  Ile Met Val Thr Phe  Arg Asn Gln
    1160                1165                1170
```

```
Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser Leu Ile  Ser Tyr Glu
    1175                1180                1185

Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg Lys Asn  Phe Val Lys
    1190                1195                1200

Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys Val Gln  His His Met
    1205                1210                1215

Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys Ala Trp  Ala Tyr Phe
    1220                1225                1230

Ser Asp  Val Asp Leu Glu Lys  Asp Val His Ser Gly  Leu Ile Gly
    1235                1240                1245

Pro Leu  Leu Val Cys His Thr  Asn Thr Leu Asn Pro  Ala His Gly
    1250                1255                1260

Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu Phe Phe  Thr Ile Phe
    1265                1270                1275

Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu Asn Met  Glu Arg Asn
    1280                1285                1290

Cys Arg  Ala Pro Cys Asn Ile  Gln Met Glu Asp Pro  Thr Phe Lys
    1295                1300                1305

Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly Tyr Ile  Met Asp Thr
    1310                1315                1320

Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln Arg Ile  Arg Trp Tyr
    1325                1330                1335

Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile His Ser  Ile His Phe
    1340                1345                1350

Ser Gly  His Val Phe Thr Val  Arg Lys Lys Glu Glu  Tyr Lys Met
    1355                1360                1365

Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe Glu Thr  Val Glu Met
    1370                1375                1380

Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val Glu Cys  Leu Ile Gly
    1385                1390                1395

Glu His  Leu His Ala Gly Met  Ser Thr Leu Phe Leu  Val Tyr Ser
    1400                1405                1410

Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala Ser Gly  His Ile Arg
    1415                1420                1425
```

```
Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr Gly Gln  Trp Ala Pro
    1430                1435                1440

Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser Ile Asn  Ala Trp Ser
    1445                1450                1455

Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val Asp Leu  Leu Ala Pro
    1460                1465                1470

Met Ile  Ile His Gly Ile Lys  Thr Gln Gly Ala Arg  Gln Lys Phe
    1475                1480                1485

Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile Met Tyr  Ser Leu Asp
    1490                1495                1500

Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn Ser Thr  Gly Thr Leu
    1505                1510                1515

Met Val  Phe Phe Gly Asn Val  Asp Ser Ser Gly Ile  Lys His Asn
    1520                1525                1530

Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr Ile Arg  Leu His Pro
    1535                1540                1545

Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg Met Glu  Leu Met Gly
    1550                1555                1560

Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu Gly Met  Glu Ser Lys
    1565                1570                1575

Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser Ser Tyr  Phe Thr Asn
    1580                1585                1590

Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala Arg Leu  His Leu Gln
    1595                1600                1605

Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val Asn Asn  Pro Lys Glu
    1610                1615                1620

Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met Lys Val  Thr Gly Val
    1625                1630                1635

Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr Ser Met  Tyr Val Lys
    1640                1645                1650

Glu Phe  Leu Ile Ser Ser Ser  Gln Asp Gly His Gln  Trp Thr Leu
    1655                1660                1665

Phe Phe  Gln Asn Gly Lys Val  Lys Val Phe Gln Gly  Asn Gln Asp
    1670                1675                1680
```

```
Ser Phe  Thr Pro Val Val Asn  Ser Leu Asp Pro Pro  Leu Leu Thr
    1685                1690                1695

Arg Tyr  Leu Arg Ile His Pro  Gln Ser Trp Val His  Gln Ile Ala
    1700                1705                1710

Leu Arg  Met Glu Val Leu Gly  Cys Glu Ala Gln Asp  Leu Tyr Lys
    1715                1720                1725

Leu Thr  Arg Ala Glu Thr Gly  Gly Ala Ser Pro Gly  Thr Ser Ser
    1730                1735                1740

Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
    1745                1750                1755

Pro Gly  Ser Ser Pro Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Thr
    1760                1765                1770

Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro
    1775                1780                1785

Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1790                1795                1800

Gly Thr  Gly Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    1805                1810                1815

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser
    1820                1825                1830

Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Thr  Pro Gly Ser
    1835                1840                1845

Gly Thr  Ala Ser Ser Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1850                1855                1860

Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    1865                1870                1875

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser
    1880                1885                1890

Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1895                1900                1905

Thr Ser  Ser Thr Gly Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala
    1910                1915                1920

Ser Ser  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1925                1930                1935
```

```
Pro Gly  Ser Ser Pro Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ser
    1940                1945                1950

Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro Gly Ser  Ser Thr Pro
    1955                1960                1965

Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1970                1975                1980

Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    1985                1990                1995

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    2000                2005                2010

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Thr  Pro Gly Ser
    2015                2020                2025

Gly Thr  Ala Ser Ser Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    2030                2035                2040

Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    2045                2050                2055

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ser
    2060                2065                2070

Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro Gly Thr  Pro Gly Ser
    2075                2080                2085

Gly Thr  Ala Ser Ser Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    2090                2095                2100

Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    2105                2110                2115

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ser
    2120                2125                2130

Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Ser  Ser Thr Pro
    2135                2140                2145

Ser Gly  Ala Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    2150                2155                2160

Thr Gly  Ser Pro Gly Thr Pro  Gly Ser Gly Thr Ala  Ser Ser Ser
    2165                2170                2175

Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr Gly Ser  Pro Gly Ser
    2180                2185                2190
```

```
Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Ser  Ser Thr Pro
    2195                 2200                 2205

Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    2210                 2215                 2220

Gly Thr  Gly Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    2225                 2230                 2235

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Thr
    2240                 2245                 2250

Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro Gly Ala  Ser Pro Gly
    2255                 2260                 2265

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    2270                 2275                 2280

Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    2285                 2290                 2295

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Thr
    2300                 2305                 2310

Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro
    2315                 2320                 2325

Ser Gly  Ala Thr Gly Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala
    2330                 2335                 2340

Ser Ser  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    2345                 2350                 2355

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser
    2360                 2365                 2370

Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Ser  Ser Thr Pro
    2375                 2380                 2385

Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    2390                 2395                 2400

Gly Thr  Gly Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
    2405                 2410                 2415

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Thr
    2420                 2425                 2430

Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro
    2435                 2440                 2445
```

```
Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    2450                 2455                 2460

Gly Thr  Gly Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
    2465                 2470                 2475

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    2480                 2485                 2490

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ser  Ser Thr Pro
    2495                 2500                 2505

Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    2510                 2515                 2520

Gly Thr  Gly Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    2525                 2530                 2535

Pro Gly  Ser Ser Pro Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Thr
    2540                 2545                 2550

Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro
    2555                 2560                 2565

Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    2570                 2575                 2580

Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    2585                 2590                 2595

Pro
```

<210> 1587
<211> 2599
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1587

```
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1               5                  10                  15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45
```

```
Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320
```

```
Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590
```

```
Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Gly Pro Gly Ala Ser Pro Gly
            740                 745                 750

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
        755                 760                 765

Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
    770                 775                 780

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser
785                 790                 795                 800

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
                805                 810                 815

Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly
            820                 825                 830

Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
        835                 840                 845

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
    850                 855                 860
```

Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
865 870 875 880

Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser
885 890 895

Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
900 905 910

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly
915 920 925

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly
930 935 940

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr
945 950 955 960

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly
965 970 975

Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly
980 985 990

Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly
995 1000 1005

Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
1010 1015 1020

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Pro Val Leu
1025 1030 1035

Lys Arg His Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp
1040 1045 1050

Gln Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys
1055 1060 1065

Lys Glu Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro
1070 1075 1080

Arg Ser Phe Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val
1085 1090 1095

Glu Arg Leu Trp Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu
1100 1105 1110

Arg Asn Arg Ala Gln Ser Gly Ser Val Pro Gln Phe Lys Lys Val
1115 1120 1125

```
Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe Thr Gln  Pro Leu Tyr
    1130                1135                1140

Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu Leu Gly  Pro Tyr Ile
    1145                1150                1155

Arg Ala  Glu Val Glu Asp Asn  Ile Met Val Thr Phe  Arg Asn Gln
    1160                1165                1170

Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser Leu Ile  Ser Tyr Glu
    1175                1180                1185

Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg Lys Asn  Phe Val Lys
    1190                1195                1200

Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys Val Gln  His His Met
    1205                1210                1215

Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys Ala Trp  Ala Tyr Phe
    1220                1225                1230

Ser Asp  Val Asp Leu Glu Lys  Asp Val His Ser Gly  Leu Ile Gly
    1235                1240                1245

Pro Leu  Leu Val Cys His Thr  Asn Thr Leu Asn Pro  Ala His Gly
    1250                1255                1260

Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu Phe Phe  Thr Ile Phe
    1265                1270                1275

Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu Asn Met  Glu Arg Asn
    1280                1285                1290

Cys Arg  Ala Pro Cys Asn Ile  Gln Met Glu Asp Pro  Thr Phe Lys
    1295                1300                1305

Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly Tyr Ile  Met Asp Thr
    1310                1315                1320

Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln Arg Ile  Arg Trp Tyr
    1325                1330                1335

Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile His Ser  Ile His Phe
    1340                1345                1350

Ser Gly  His Val Phe Thr Val  Arg Lys Lys Glu Glu  Tyr Lys Met
    1355                1360                1365

Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe Glu Thr  Val Glu Met
    1370                1375                1380
```

```
Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val Glu Cys  Leu Ile Gly
    1385                1390                1395

Glu His  Leu His Ala Gly Met  Ser Thr Leu Phe Leu  Val Tyr Ser
    1400                1405                1410

Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala Ser Gly  His Ile Arg
    1415                1420                1425

Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr Gly Gln  Trp Ala Pro
    1430                1435                1440

Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser Ile Asn  Ala Trp Ser
    1445                1450                1455

Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val Asp Leu  Leu Ala Pro
    1460                1465                1470

Met Ile  Ile His Gly Ile Lys  Thr Gln Gly Ala Arg  Gln Lys Phe
    1475                1480                1485

Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile Met Tyr  Ser Leu Asp
    1490                1495                1500

Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn Ser Thr  Gly Thr Leu
    1505                1510                1515

Met Val  Phe Phe Gly Asn Val  Asp Ser Ser Gly Ile  Lys His Asn
    1520                1525                1530

Ile Phe  Asn Pro Pro Ile Ile  Ala Arg Tyr Ile Arg  Leu His Pro
    1535                1540                1545

Thr His  Tyr Ser Ile Arg Ser  Thr Leu Arg Met Glu  Leu Met Gly
    1550                1555                1560

Cys Asp  Leu Asn Ser Cys Ser  Met Pro Leu Gly Met  Glu Ser Lys
    1565                1570                1575

Ala Ile  Ser Asp Ala Gln Ile  Thr Ala Ser Ser Tyr  Phe Thr Asn
    1580                1585                1590

Met Phe  Ala Thr Trp Ser Pro  Ser Lys Ala Arg Leu  His Leu Gln
    1595                1600                1605

Gly Arg  Ser Asn Ala Trp Arg  Pro Gln Val Asn Asn  Pro Lys Glu
    1610                1615                1620

Trp Leu  Gln Val Asp Phe Gln  Lys Thr Met Lys Val  Thr Gly Val
    1625                1630                1635
```

```
Thr Thr  Gln Gly Val Lys Ser  Leu Leu Thr Ser Met  Tyr Val Lys
    1640                1645                1650

Glu Phe  Leu Ile Ser Ser Ser  Gln Asp Gly His Gln  Trp Thr Leu
    1655                1660                1665

Phe Phe  Gln Asn Gly Lys Val  Lys Val Phe Gln Gly  Asn Gln Asp
    1670                1675                1680

Ser Phe  Thr Pro Val Val Asn  Ser Leu Asp Pro Pro  Leu Leu Thr
    1685                1690                1695

Arg Tyr  Leu Arg Ile His Pro  Gln Ser Trp Val His  Gln Ile Ala
    1700                1705                1710

Leu Arg  Met Glu Val Leu Gly  Cys Glu Ala Gln Asp  Leu Tyr Lys
    1715                1720                1725

Leu Thr  Arg Ala Glu Thr Gly  Gly Ala Ser Pro Gly  Thr Ser Ser
    1730                1735                1740

Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala Ser Thr  Gly Thr Gly
    1745                1750                1755

Pro Gly  Ser Ser Pro Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Thr
    1760                1765                1770

Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro Gly Ser  Ser Thr Pro
    1775                1780                1785

Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser Pro Ser  Ala Ser Thr
    1790                1795                1800

Gly Thr  Gly Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    1805                1810                1815

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser
    1820                1825                1830

Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Thr  Pro Gly Ser
    1835                1840                1845

Gly Thr  Ala Ser Ser Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1850                1855                1860

Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    1865                1870                1875

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser
    1880                1885                1890
```

```
Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1895                1900                1905

Thr Ser  Ser Thr Gly Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala
    1910                1915                1920

Ser Ser  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1925                1930                1935

Pro Gly  Ser Ser Pro Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ser
    1940                1945                1950

Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro Gly Ser  Ser Thr Pro
    1955                1960                1965

Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1970                1975                1980

Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    1985                1990                1995

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ala
    2000                2005                2010

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Thr  Pro Gly Ser
    2015                2020                2025

Gly Thr  Ala Ser Ser Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    2030                2035                2040

Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    2045                2050                2055

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ser
    2060                2065                2070

Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro Gly Thr  Pro Gly Ser
    2075                2080                2085

Gly Thr  Ala Ser Ser Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    2090                2095                2100

Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr Ser Ser  Thr Gly Ser
    2105                2110                2115

Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr Gly Ser  Pro Gly Ser
    2120                2125                2130

Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Ser  Ser Thr Pro
    2135                2140                2145
```

Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
2150 2155 2160

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
2165 2170 2175

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser
2180 2185 2190

Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
2195 2200 2205

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
2210 2215 2220

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
2225 2230 2235

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr
2240 2245 2250

Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly
2255 2260 2265

Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
2270 2275 2280

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
2285 2290 2295

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr
2300 2305 2310

Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
2315 2320 2325

Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
2330 2335 2340

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
2345 2350 2355

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser
2360 2365 2370

Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro
2375 2380 2385

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
2390 2395 2400

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
2405 2410 2415

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr
2420 2425 2430

Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
2435 2440 2445

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
2450 2455 2460

Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
2465 2470 2475

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala
2480 2485 2490

Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro
2495 2500 2505

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr
2510 2515 2520

Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
2525 2530 2535

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr
2540 2545 2550

Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro
2555 2560 2565

Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
2570 2575 2580

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
2585 2590 2595

Pro

<210> 1588
<211> 1763
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1588
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

```
Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285
```

```
Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560
```

```
Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830
```

```
Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1025                1030                1035

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1040                1045                1050

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1055                1060                1065

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1070                1075                1080

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1085                1090                1095
```

```
Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1100                1105                1110

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1115                1120                1125

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1130                1135                1140

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1145                1150                1155

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1160                1165                1170

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1175                1180                1185

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1190                1195                1200

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1205                1210                1215

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1220                1225                1230

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1235                1240                1245

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1250                1255                1260

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1265                1270                1275

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1280                1285                1290

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1295                1300                1305

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1310                1315                1320

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1325                1330                1335

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1340                1345                1350
```

```
Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1355                 1360                 1365

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1370                 1375                 1380

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1385                 1390                 1395

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1400                 1405                 1410

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1415                 1420                 1425

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Lys Leu  Thr Arg Ala
    1430                 1435                 1440

Glu Thr  Gly Ala Gly Ser Pro  Gly Ala Glu Thr Ala  Pro Gly Ala
    1445                 1450                 1455

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ala  Ser Pro Gly
    1460                 1465                 1470

Thr Ser  Ser Thr Gly Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala
    1475                 1480                 1485

Ser Ser  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1490                 1495                 1500

Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser Ser Ser  Pro Gly Ser
    1505                 1510                 1515

Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro Gly Thr  Pro Gly Ser
    1520                 1525                 1530

Gly Thr  Ala Ser Ser Ser Pro  Gly Ser Ser Thr Pro  Ser Gly Ala
    1535                 1540                 1545

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1550                 1555                 1560

Pro Gly  Ser Ser Pro Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ser
    1565                 1570                 1575

Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro Gly Ala  Ser Pro Gly
    1580                 1585                 1590

Thr Ser  Ser Thr Gly Ser Pro  Gly Thr Pro Gly Ser  Gly Thr Ala
    1595                 1600                 1605
```

```
Ser Ser  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1610                 1615                 1620

Pro Gly  Ser Ser Pro Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ser
    1625                 1630                 1635

Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro Gly Ala  Ser Pro Gly
    1640                 1645                 1650

Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser Pro Gly  Thr Ser Ser
    1655                 1660                 1665

Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1670                 1675                 1680

Pro Gly  Ser Ser Pro Ser Ala  Ser Thr Gly Thr Gly  Pro Gly Ala
    1685                 1690                 1695

Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro Gly Ser  Ser Pro Ser
    1700                 1705                 1710

Ala Ser  Thr Gly Thr Gly Pro  Gly Thr Pro Gly Ser  Gly Thr Ala
    1715                 1720                 1725

Ser Ser  Ser Pro Gly Ser Ser  Thr Pro Ser Gly Ala  Thr Gly Ser
    1730                 1735                 1740

Gly Ala  Glu Thr Ala Glu Gln  Lys Leu Ile Ser Glu  Glu Asp Leu
    1745                 1750                 1755

Ser Pro  Ala Thr Gly
    1760


<210> 1589
<211> 2310
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
      polypeptide

<400> 1589
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1               5                  10                  15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45
```

```
Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320
```

```
Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590
```

```
Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860
```

```
Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1025                1030                1035

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1040                1045                1050

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1055                1060                1065

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1070                1075                1080

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1085                1090                1095

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1100                1105                1110

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1115                1120                1125
```

```
Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1130                1135                1140

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1145                1150                1155

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1160                1165                1170

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1175                1180                1185

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1190                1195                1200

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1205                1210                1215

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1220                1225                1230

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1235                1240                1245

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1250                1255                1260

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1265                1270                1275

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1280                1285                1290

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1295                1300                1305

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1310                1315                1320

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1325                1330                1335

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1340                1345                1350

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1355                1360                1365

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1370                1375                1380
```

```
Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1385                1390                1395

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1400                1405                1410

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1415                1420                1425

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Lys Leu  Thr Arg Ala
    1430                1435                1440

Glu Thr  Gly Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1445                1450                1455

Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr Gly Pro  Gly Ser Ser
    1460                1465                1470

Pro Ser  Ala Ser Thr Gly Thr  Gly Pro Gly Thr Pro  Gly Ser Gly
    1475                1480                1485

Thr Ala  Ser Ser Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    1490                1495                1500

Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro
    1505                1510                1515

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Thr Pro
    1520                1525                1530

Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly Ser Ser  Thr Pro Ser
    1535                1540                1545

Gly Ala  Thr Gly Ser Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    1550                1555                1560

Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1565                1570                1575

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Thr Pro
    1580                1585                1590

Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly Ser Ser  Thr Pro Ser
    1595                1600                1605

Gly Ala  Thr Gly Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
    1610                1615                1620

Gly Ser  Pro Gly Thr Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro
    1625                1630                1635
```

```
Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Gly Ser Ser
    1640                1645                1650

Pro Ser  Ala Ser Thr Gly Thr  Gly Pro Gly Ser Ser  Pro Ser Ala
    1655                1660                1665

Ser Thr  Gly Thr Gly Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    1670                1675                1680

Gly Ser  Pro Gly Ser Ser Thr  Pro Ser Gly Ala Thr  Gly Ser Pro
    1685                1690                1695

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1700                1705                1710

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ala Ser  Pro Gly Thr
    1715                1720                1725

Ser Ser  Thr Gly Ser Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    1730                1735                1740

Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1745                1750                1755

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1760                1765                1770

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Pro Ser Ala
    1775                1780                1785

Ser Thr  Gly Thr Gly Pro Gly  Thr Pro Gly Ser Gly  Thr Ala Ser
    1790                1795                1800

Ser Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1805                1810                1815

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1820                1825                1830

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser
    1835                1840                1845

Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    1850                1855                1860

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1865                1870                1875

Gly Thr  Pro Gly Ser Gly Thr  Ala Ser Ser Ser Pro  Gly Ser Ser
    1880                1885                1890
```

```
Thr Pro  Ser Gly Ala Thr Gly  Ser Pro Gly Ser Ser  Thr Pro Ser
    1895                1900                1905

Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    1910                1915                1920

Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro
    1925                1930                1935

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    1940                1945                1950

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Thr Pro  Gly Ser Gly
    1955                1960                1965

Thr Ala  Ser Ser Ser Pro Gly  Ala Ser Pro Gly Thr  Ser Ser Thr
    1970                1975                1980

Gly Ser  Pro Gly Ala Ser Pro  Gly Thr Ser Ser Thr  Gly Ser Pro
    1985                1990                1995

Gly Ala  Ser Pro Gly Thr Ser  Ser Thr Gly Ser Pro  Gly Ala Ser
    2000                2005                2010

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Thr Pro  Gly Ser Gly
    2015                2020                2025

Thr Ala  Ser Ser Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    2030                2035                2040

Gly Ser  Pro Gly Thr Pro Gly  Ser Gly Thr Ala Ser  Ser Ser Pro
    2045                2050                2055

Gly Ser  Ser Thr Pro Ser Gly  Ala Thr Gly Ser Pro  Gly Thr Pro
    2060                2065                2070

Gly Ser  Gly Thr Ala Ser Ser  Ser Pro Gly Ser Ser  Thr Pro Ser
    2075                2080                2085

Gly Ala  Thr Gly Ser Pro Gly  Ser Ser Thr Pro Ser  Gly Ala Thr
    2090                2095                2100

Gly Ser  Pro Gly Ser Ser Pro  Ser Ala Ser Thr Gly  Thr Gly Pro
    2105                2110                2115

Gly Ser  Ser Pro Ser Ala Ser  Thr Gly Thr Gly Pro  Gly Ala Ser
    2120                2125                2130

Pro Gly  Thr Ser Ser Thr Gly  Ser Pro Gly Thr Pro  Gly Ser Gly
    2135                2140                2145
```

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
2150 2155 2160

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
2165 2170 2175

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
2180 2185 2190

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr
2195 2200 2205

Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
2210 2215 2220

Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
2225 2230 2235

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
2240 2245 2250

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly
2255 2260 2265

Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr
2270 2275 2280

Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
2285 2290 2295

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
2300 2305 2310

<210> 1590
<211> 2310
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1590
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
50 55 60

```
Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335
```

```
Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605
```

```
Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Ala Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880
```

```
Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
               885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
           900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
       915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
   930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
               965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
           980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
       995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
   1010                1015                1020

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
   1025                1030                1035

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
   1040                1045                1050

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
   1055                1060                1065

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
   1070                1075                1080

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
   1085                1090                1095

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
   1100                1105                1110

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
   1115                1120                1125

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
   1130                1135                1140
```

```
Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1145                1150                1155

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1160                1165                1170

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1175                1180                1185

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1190                1195                1200

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1205                1210                1215

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1220                1225                1230

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1235                1240                1245

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1250                1255                1260

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1265                1270                1275

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1280                1285                1290

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1295                1300                1305

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1310                1315                1320

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1325                1330                1335

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1340                1345                1350

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1355                1360                1365

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1370                1375                1380

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1385                1390                1395
```

```
Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1400                1405                1410

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1415                1420                1425

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr Lys Leu  Thr Arg Ala
    1430                1435                1440

Glu Thr  Gly Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1445                1450                1455

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1460                1465                1470

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1475                1480                1485

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1490                1495                1500

Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1505                1510                1515

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    1520                1525                1530

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Ser Glu  Pro Ala Thr
    1535                1540                1545

Ser Gly  Ser Glu Thr Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1550                1555                1560

Thr Glu  Glu Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1565                1570                1575

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1580                1585                1590

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Ser Pro  Ala Gly Ser
    1595                1600                1605

Pro Thr  Ser Thr Glu Glu Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1610                1615                1620

Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1625                1630                1635

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    1640                1645                1650
```

```
Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala
    1655                 1660                 1665

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1670                 1675                 1680

Glu Thr  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1685                 1690                 1695

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1700                 1705                 1710

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Thr Ser  Glu Ser Ala
    1715                 1720                 1725

Thr Pro  Glu Ser Gly Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1730                 1735                 1740

Thr Glu  Glu Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1745                 1750                 1755

Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser
    1760                 1765                 1770

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro
    1775                 1780                 1785

Ser Glu  Gly Ser Ala Pro Gly  Thr Ser Thr Glu Pro  Ser Glu Gly
    1790                 1795                 1800

Ser Ala  Pro Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1805                 1810                 1815

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1820                 1825                 1830

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Thr Glu Pro
    1835                 1840                 1845

Ser Glu  Gly Ser Ala Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1850                 1855                 1860

Thr Glu  Glu Gly Thr Ser Thr  Glu Pro Ser Glu Gly  Ser Ala Pro
    1865                 1870                 1875

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Ser Glu
    1880                 1885                 1890

Pro Ala  Thr Ser Gly Ser Glu  Thr Pro Gly Thr Ser  Glu Ser Ala
    1895                 1900                 1905
```

```
Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    1910                 1915                 1920

Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    1925                 1930                 1935

Gly Thr  Ser Thr Glu Pro Ser  Glu Gly Ser Ala Pro  Gly Thr Ser
    1940                 1945                 1950

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Pro  Ala Gly Ser
    1955                 1960                 1965

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    1970                 1975                 1980

Thr Glu  Glu Gly Ser Pro Ala  Gly Ser Pro Thr Ser  Thr Glu Glu
    1985                 1990                 1995

Gly Thr  Ser Glu Ser Ala Thr  Pro Glu Ser Gly Pro  Gly Thr Ser
    2000                 2005                 2010

Thr Glu  Pro Ser Glu Gly Ser  Ala Pro Gly Thr Ser  Glu Ser Ala
    2015                 2020                 2025

Thr Pro  Glu Ser Gly Pro Gly  Ser Glu Pro Ala Thr  Ser Gly Ser
    2030                 2035                 2040

Glu Thr  Pro Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    2045                 2050                 2055

Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser
    2060                 2065                 2070

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Thr Ser  Thr Glu Pro
    2075                 2080                 2085

Ser Glu  Gly Ser Ala Pro Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    2090                 2095                 2100

Thr Glu  Glu Gly Thr Ser Glu  Ser Ala Thr Pro Glu  Ser Gly Pro
    2105                 2110                 2115

Gly Ser  Glu Pro Ala Thr Ser  Gly Ser Glu Thr Pro  Gly Thr Ser
    2120                 2125                 2130

Glu Ser  Ala Thr Pro Glu Ser  Gly Pro Gly Ser Pro  Ala Gly Ser
    2135                 2140                 2145

Pro Thr  Ser Thr Glu Glu Gly  Ser Pro Ala Gly Ser  Pro Thr Ser
    2150                 2155                 2160
```

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
2165 2170 2175

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser
2180 2185 2190

Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala
2195 2200 2205

Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
2210 2215 2220

Glu Thr Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
2225 2230 2235

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser
2240 2245 2250

Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro
2255 2260 2265

Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
2270 2275 2280

Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
2285 2290 2295

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
2300 2305 2310

<210> 1591
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 1591
Ala Ser Thr Gly Glu Thr Gly Glu Thr Gly Glu
1 5 10

<210> 1592
<211> 2332
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1592
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

```
Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285
```

```
Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560
```

```
Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Ser Arg His Pro Ser Thr Arg
            740                 745                 750

Gln Lys Gln Phe Asn Ala Thr Thr Ile Pro Glu Asn Asp Ile Glu Lys
        755                 760                 765

Thr Asp Pro Trp Phe Ala His Arg Thr Pro Met Pro Lys Ile Gln Asn
    770                 775                 780

Val Ser Ser Ser Asp Leu Leu Met Leu Leu Arg Gln Ser Pro Thr Pro
785                 790                 795                 800

His Gly Leu Ser Leu Ser Asp Leu Gln Glu Ala Lys Tyr Glu Thr Phe
                805                 810                 815

Ser Asp Asp Pro Ser Pro Gly Ala Ile Asp Ser Asn Asn Ser Leu Ser
            820                 825                 830
```

```
Glu Met Thr His Phe Arg Pro Gln Leu His His Ser Gly Asp Met Val
        835                 840                 845

Phe Thr Pro Glu Ser Gly Leu Gln Leu Arg Leu Asn Glu Lys Leu Gly
    850                 855                 860

Thr Thr Ala Ala Thr Glu Leu Lys Lys Leu Asp Phe Lys Val Ser Ser
865                 870                 875                 880

Thr Ser Asn Asn Leu Ile Ser Thr Ile Pro Ser Asp Asn Leu Ala Ala
                885                 890                 895

Gly Thr Asp Asn Thr Ser Ser Leu Gly Pro Pro Ser Met Pro Val His
            900                 905                 910

Tyr Asp Ser Gln Leu Asp Thr Thr Leu Phe Gly Lys Lys Ser Ser Pro
        915                 920                 925

Leu Thr Glu Ser Gly Gly Pro Leu Ser Leu Ser Glu Glu Asn Asn Asp
    930                 935                 940

Ser Lys Leu Leu Glu Ser Gly Leu Met Asn Ser Gln Glu Ser Ser Trp
945                 950                 955                 960

Gly Lys Asn Val Ser Ser Thr Glu Ser Gly Arg Leu Phe Lys Gly Lys
                965                 970                 975

Arg Ala His Gly Pro Ala Leu Leu Thr Lys Asp Asn Ala Leu Phe Lys
            980                 985                 990

Val Ser Ile Ser Leu Leu Lys Thr  Asn Lys Thr Ser Asn  Asn Ser Ala
        995                 1000                1005

Thr Asn  Arg Lys Thr His Ile  Asp Gly Pro Ser Leu  Leu Ile Glu
    1010                1015                1020

Asn Ser  Pro Ser Val Trp Gln  Asn Ile Leu Glu Ser  Asp Thr Glu
    1025                1030                1035

Phe Lys  Lys Val Thr Pro Leu  Ile His Asp Arg Met  Leu Met Asp
    1040                1045                1050

Lys Asn  Ala Thr Ala Leu Arg  Leu Asn His Met Ser  Asn Lys Thr
    1055                1060                1065

Thr Ser  Ser Lys Asn Met Glu  Met Val Gln Gln Lys  Lys Glu Gly
    1070                1075                1080

Pro Ile  Pro Pro Asp Ala Gln  Asn Pro Asp Met Ser  Phe Phe Lys
    1085                1090                1095
```

```
Met Leu  Phe Leu Pro Glu Ser  Ala Arg Trp Ile Gln  Arg Thr His
    1100                1105                1110

Gly Lys  Asn Ser Leu Asn Ser  Gly Gln Gly Pro Ser  Pro Lys Gln
    1115                1120                1125

Leu Val  Ser Leu Gly Pro Glu  Lys Ser Val Glu Gly  Gln Asn Phe
    1130                1135                1140

Leu Ser  Glu Lys Asn Lys Val  Val Val Gly Lys Gly  Glu Phe Thr
    1145                1150                1155

Lys Asp  Val Gly Leu Lys Glu  Met Val Phe Pro Ser  Ser Arg Asn
    1160                1165                1170

Leu Phe  Leu Thr Asn Leu Asp  Asn Leu His Glu Asn  Asn Thr His
    1175                1180                1185

Asn Gln  Glu Lys Lys Ile Gln  Glu Glu Ile Glu Lys  Lys Glu Thr
    1190                1195                1200

Leu Ile  Gln Glu Asn Val Val  Leu Pro Gln Ile His  Thr Val Thr
    1205                1210                1215

Gly Thr  Lys Asn Phe Met Lys  Asn Leu Phe Leu Leu  Ser Thr Arg
    1220                1225                1230

Gln Asn  Val Glu Gly Ser Tyr  Asp Gly Ala Tyr Ala  Pro Val Leu
    1235                1240                1245

Gln Asp  Phe Arg Ser Leu Asn  Asp Ser Thr Asn Arg  Thr Lys Lys
    1250                1255                1260

His Thr  Ala His Phe Ser Lys  Lys Gly Glu Glu Glu  Asn Leu Glu
    1265                1270                1275

Gly Leu  Gly Asn Gln Thr Lys  Gln Ile Val Glu Lys  Tyr Ala Cys
    1280                1285                1290

Thr Thr  Arg Ile Ser Pro Asn  Thr Ser Gln Gln Asn  Phe Val Thr
    1295                1300                1305

Gln Arg  Ser Lys Arg Ala Leu  Lys Gln Phe Arg Leu  Pro Leu Glu
    1310                1315                1320

Glu Thr  Glu Leu Glu Lys Arg  Ile Ile Val Asp Asp  Thr Ser Thr
    1325                1330                1335

Gln Trp  Ser Lys Asn Met Lys  His Leu Thr Pro Ser  Thr Leu Thr
    1340                1345                1350
```

```
Gln Ile  Asp Tyr Asn Glu Lys  Glu Lys Gly Ala Ile  Thr Gln Ser
    1355                 1360                 1365

Pro Leu  Ser Asp Cys Leu Thr  Arg Ser His Ser Ile  Pro Gln Ala
    1370                 1375                 1380

Asn Arg  Ser Pro Leu Pro Ile  Ala Lys Val Ser Ser  Phe Pro Ser
    1385                 1390                 1395

Ile Arg  Pro Ile Tyr Leu Thr  Arg Val Leu Phe Gln  Asp Asn Ser
    1400                 1405                 1410

Ser His  Leu Pro Ala Ala Ser  Tyr Arg Lys Lys Asp  Ser Gly Val
    1415                 1420                 1425

Gln Glu  Ser Ser His Phe Leu  Gln Gly Ala Lys Lys  Asn Asn Leu
    1430                 1435                 1440

Ser Leu  Ala Ile Leu Thr Leu  Glu Met Thr Gly Asp  Gln Arg Glu
    1445                 1450                 1455

Val Gly  Ser Leu Gly Thr Ser  Ala Thr Asn Ser Val  Thr Tyr Lys
    1460                 1465                 1470

Lys Val  Glu Asn Thr Val Leu  Pro Lys Pro Asp Leu  Pro Lys Thr
    1475                 1480                 1485

Ser Gly  Lys Val Glu Leu Leu  Pro Lys Val His Ile  Tyr Gln Lys
    1490                 1495                 1500

Asp Leu  Phe Pro Thr Glu Thr  Ser Asn Gly Ser Pro  Gly His Leu
    1505                 1510                 1515

Asp Leu  Val Glu Gly Ser Leu  Leu Gln Gly Thr Glu  Gly Ala Ile
    1520                 1525                 1530

Lys Trp  Asn Glu Ala Asn Arg  Pro Gly Lys Val Pro  Phe Leu Arg
    1535                 1540                 1545

Val Ala  Thr Glu Ser Ser Ala  Lys Thr Pro Ser Lys  Leu Leu Asp
    1550                 1555                 1560

Pro Leu  Ala Trp Asp Asn His  Tyr Gly Thr Gln Ile  Pro Lys Glu
    1565                 1570                 1575

Glu Trp  Lys Ser Gln Glu Lys  Ser Pro Glu Lys Thr  Ala Phe Lys
    1580                 1585                 1590

Lys Lys  Asp Thr Ile Leu Ser  Leu Asn Ala Cys Glu  Ser Asn His
    1595                 1600                 1605
```

```
Ala Ile  Ala Ala Ile Asn Glu  Gly Gln Asn Lys Pro  Glu Ile Glu
    1610                1615                1620

Val Thr  Trp Ala Lys Gln Gly  Arg Thr Glu Arg Leu  Cys Ser Gln
    1625                1630                1635

Asn Pro  Pro Val Leu Lys Arg  His Gln Arg Glu Ile  Thr Arg Thr
    1640                1645                1650

Thr Leu  Gln Ser Asp Gln Glu  Glu Ile Asp Tyr Asp  Asp Thr Ile
    1655                1660                1665

Ser Val  Glu Met Lys Lys Glu  Asp Phe Asp Ile Tyr  Asp Glu Asp
    1670                1675                1680

Glu Asn  Gln Ser Pro Arg Ser  Phe Gln Lys Lys Thr  Arg His Tyr
    1685                1690                1695

Phe Ile  Ala Ala Val Glu Arg  Leu Trp Asp Tyr Gly  Met Ser Ser
    1700                1705                1710

Ser Pro  His Val Leu Arg Asn  Arg Ala Gln Ser Gly  Ser Val Pro
    1715                1720                1725

Gln Phe  Lys Lys Val Val Phe  Gln Glu Phe Thr Asp  Gly Ser Phe
    1730                1735                1740

Thr Gln  Pro Leu Tyr Arg Gly  Glu Leu Asn Glu His  Leu Gly Leu
    1745                1750                1755

Leu Gly  Pro Tyr Ile Arg Ala  Glu Val Glu Asp Asn  Ile Met Val
    1760                1765                1770

Thr Phe  Arg Asn Gln Ala Ser  Arg Pro Tyr Ser Phe  Tyr Ser Ser
    1775                1780                1785

Leu Ile  Ser Tyr Glu Glu Asp  Gln Arg Gln Gly Ala  Glu Pro Arg
    1790                1795                1800

Lys Asn  Phe Val Lys Pro Asn  Glu Thr Lys Thr Tyr  Phe Trp Lys
    1805                1810                1815

Val Gln  His His Met Ala Pro  Thr Lys Asp Glu Phe  Asp Cys Lys
    1820                1825                1830

Ala Trp  Ala Tyr Phe Ser Asp  Val Asp Leu Glu Lys  Asp Val His
    1835                1840                1845

Ser Gly  Leu Ile Gly Pro Leu  Leu Val Cys His Thr  Asn Thr Leu
    1850                1855                1860
```

```
Asn Pro  Ala His Gly Arg Gln  Val Thr Val Gln Glu  Phe Ala Leu
    1865                 1870                 1875

Phe Phe  Thr Ile Phe Asp Glu  Thr Lys Ser Trp Tyr  Phe Thr Glu
    1880                 1885                 1890

Asn Met  Glu Arg Asn Cys Arg  Ala Pro Cys Asn Ile  Gln Met Glu
    1895                 1900                 1905

Asp Pro  Thr Phe Lys Glu Asn  Tyr Arg Phe His Ala  Ile Asn Gly
    1910                 1915                 1920

Tyr Ile  Met Asp Thr Leu Pro  Gly Leu Val Met Ala  Gln Asp Gln
    1925                 1930                 1935

Arg Ile  Arg Trp Tyr Leu Leu  Ser Met Gly Ser Asn  Glu Asn Ile
    1940                 1945                 1950

His Ser  Ile His Phe Ser Gly  His Val Phe Thr Val  Arg Lys Lys
    1955                 1960                 1965

Glu Glu  Tyr Lys Met Ala Leu  Tyr Asn Leu Tyr Pro  Gly Val Phe
    1970                 1975                 1980

Glu Thr  Val Glu Met Leu Pro  Ser Lys Ala Gly Ile  Trp Arg Val
    1985                 1990                 1995

Glu Cys  Leu Ile Gly Glu His  Leu His Ala Gly Met  Ser Thr Leu
    2000                 2005                 2010

Phe Leu  Val Tyr Ser Asn Lys  Cys Gln Thr Pro Leu  Gly Met Ala
    2015                 2020                 2025

Ser Gly  His Ile Arg Asp Phe  Gln Ile Thr Ala Ser  Gly Gln Tyr
    2030                 2035                 2040

Gly Gln  Trp Ala Pro Lys Leu  Ala Arg Leu His Tyr  Ser Gly Ser
    2045                 2050                 2055

Ile Asn  Ala Trp Ser Thr Lys  Glu Pro Phe Ser Trp  Ile Lys Val
    2060                 2065                 2070

Asp Leu  Leu Ala Pro Met Ile  Ile His Gly Ile Lys  Thr Gln Gly
    2075                 2080                 2085

Ala Arg  Gln Lys Phe Ser Ser  Leu Tyr Ile Ser Gln  Phe Ile Ile
    2090                 2095                 2100

Met Tyr  Ser Leu Asp Gly Lys  Lys Trp Gln Thr Tyr  Arg Gly Asn
    2105                 2110                 2115
```

Ser Thr Gly Thr Leu Met Val Phe Phe Gly Asn Val Asp Ser Ser
2120 2125 2130

Gly Ile Lys His Asn Ile Phe Asn Pro Pro Ile Ile Ala Arg Tyr
2135 2140 2145

Ile Arg Leu His Pro Thr His Tyr Ser Ile Arg Ser Thr Leu Arg
2150 2155 2160

Met Glu Leu Met Gly Cys Asp Leu Asn Ser Cys Ser Met Pro Leu
2165 2170 2175

Gly Met Glu Ser Lys Ala Ile Ser Asp Ala Gln Ile Thr Ala Ser
2180 2185 2190

Ser Tyr Phe Thr Asn Met Phe Ala Thr Trp Ser Pro Ser Lys Ala
2195 2200 2205

Arg Leu His Leu Gln Gly Arg Ser Asn Ala Trp Arg Pro Gln Val
2210 2215 2220

Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Gln Lys Thr Met
2225 2230 2235

Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser Leu Leu Thr
2240 2245 2250

Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly
2255 2260 2265

His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val Lys Val Phe
2270 2275 2280

Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn Ser Leu Asp
2285 2290 2295

Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro Gln Ser Trp
2300 2305 2310

Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly Cys Glu Ala
2315 2320 2325

Gln Asp Leu Tyr
2330

<210> 1593
<211> 1438
<212> PRT
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1593

```
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1               5                  10                  15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
            20                  25                  30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
        35                  40                  45

Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro
    50                  55                  60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65                  70                  75                  80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
                85                  90                  95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
            100                 105                 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255
```

```
Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525
```

```
Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800
```

```
Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1025                1030                1035

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1040                1045                1050

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1055                1060                1065
```

```
Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1070                1075                1080

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1085                1090                1095

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1100                1105                1110

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1115                1120                1125

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1130                1135                1140

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1145                1150                1155

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1160                1165                1170

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1175                1180                1185

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1190                1195                1200

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1205                1210                1215

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1220                1225                1230

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1235                1240                1245

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1250                1255                1260

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1265                1270                1275

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1280                1285                1290

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1295                1300                1305

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1310                1315                1320
```

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1325 1330 1335

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1340 1345 1350

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1355 1360 1365

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1370 1375 1380

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1385 1390 1395

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1400 1405 1410

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1415 1420 1425

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr
1430 1435

<210> 1594
<211> 1438
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1594
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

```
Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
        115                 120                 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
    130                 135                 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145                 150                 155                 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
                165                 170                 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
            180                 185                 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
        195                 200                 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
    210                 215                 220

Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380
```

```
Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655
```

```
Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765

Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925
```

```
Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1025                1030                1035

Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1040                1045                1050

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1055                1060                1065

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1070                1075                1080

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1085                1090                1095

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1100                1105                1110

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1115                1120                1125

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1130                1135                1140

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1145                1150                1155

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1160                1165                1170

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1175                1180                1185
```

```
Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1190                1195                1200

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1205                1210                1215

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1220                1225                1230

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1235                1240                1245

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1250                1255                1260

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1265                1270                1275

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1280                1285                1290

Ala Gln  Ile Thr Ala Ser Ser  Tyr Phe Thr Asn Met  Phe Ala Thr
    1295                1300                1305

Trp Ser  Pro Ser Lys Ala Arg  Leu His Leu Gln Gly  Arg Ser Asn
    1310                1315                1320

Ala Trp  Arg Pro Gln Val Asn  Asn Pro Lys Glu Trp  Leu Gln Val
    1325                1330                1335

Asp Phe  Gln Lys Thr Met Lys  Val Thr Gly Val Thr  Thr Gln Gly
    1340                1345                1350

Val Lys  Ser Leu Leu Thr Ser  Met Tyr Val Lys Glu  Phe Leu Ile
    1355                1360                1365

Ser Ser  Ser Gln Asp Gly His  Gln Trp Thr Leu Phe  Phe Gln Asn
    1370                1375                1380

Gly Lys  Val Lys Val Phe Gln  Gly Asn Gln Asp Ser  Phe Thr Pro
    1385                1390                1395

Val Val  Asn Ser Leu Asp Pro  Pro Leu Leu Thr Arg  Tyr Leu Arg
    1400                1405                1410

Ile His  Pro Gln Ser Trp Val  His Gln Ile Ala Leu  Arg Met Glu
    1415                1420                1425

Val Leu  Gly Cys Glu Ala Gln  Asp Leu Tyr
    1430                1435
```

<210> 1595
<211> 1438
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1595
Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser Trp Asp Tyr
1 5 10 15

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys
35 40 45

Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn
130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser
145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu
165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu
180 185 190

His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp
195 200 205

His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser
210 215 220

```
Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg
225                 230                 235                 240

Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His
                245                 250                 255

Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu
            260                 265                 270

Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile
        275                 280                 285

Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly
    290                 295                 300

Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met
305                 310                 315                 320

Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg
                325                 330                 335

Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp
            340                 345                 350

Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe
        355                 360                 365

Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His
    370                 375                 380

Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu
385                 390                 395                 400

Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro
                405                 410                 415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr
            420                 425                 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile
        435                 440                 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile
    450                 455                 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile
465                 470                 475                 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys
                485                 490                 495
```

```
His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys
            500                 505                 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys
        515                 520                 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala
    530                 535                 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp
545                 550                 555                 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe
                565                 570                 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln
            580                 585                 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe
        595                 600                 605

Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val Phe Asp Ser
    610                 615                 620

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu
625                 630                 635                 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr
                645                 650                 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro
            660                 665                 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp
        675                 680                 685

Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly Met Thr Ala
    690                 695                 700

Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp Tyr Tyr Glu
705                 710                 715                 720

Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys Asn Asn Ala
                725                 730                 735

Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His
            740                 745                 750

Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln Glu Glu Ile
        755                 760                 765
```

```
Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu Asp Phe Asp
    770                 775                 780

Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe Gln Lys Lys
785                 790                 795                 800

Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp Asp Tyr Gly
                805                 810                 815

Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln Ser Gly Ser
            820                 825                 830

Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr Asp Gly Ser
        835                 840                 845

Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His Leu Gly Leu
    850                 855                 860

Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile Met Val Thr
865                 870                 875                 880

Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser Ser Leu Ile
                885                 890                 895

Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg Lys Asn Phe
            900                 905                 910

Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val Gln His His
        915                 920                 925

Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp Ala Tyr Phe
    930                 935                 940

Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu Ile Gly Pro
945                 950                 955                 960

Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln
                965                 970                 975

Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr
            980                 985                 990

Lys Ser Trp Tyr Phe Thr Glu Asn  Met Glu Arg Asn Cys  Arg Ala Pro
        995                 1000                1005

Cys Asn  Ile Gln Met Glu Asp  Pro Thr Phe Lys Glu  Asn Tyr Arg
    1010                1015                1020

Phe His  Ala Ile Asn Gly Tyr  Ile Met Asp Thr Leu  Pro Gly Leu
    1025                1030                1035
```

```
Val Met  Ala Gln Asp Gln Arg  Ile Arg Trp Tyr Leu  Leu Ser Met
    1040                1045                1050

Gly Ser  Asn Glu Asn Ile His  Ser Ile His Phe Ser  Gly His Val
    1055                1060                1065

Phe Thr  Val Arg Lys Lys Glu  Glu Tyr Lys Met Ala  Leu Tyr Asn
    1070                1075                1080

Leu Tyr  Pro Gly Val Phe Glu  Thr Val Glu Met Leu  Pro Ser Lys
    1085                1090                1095

Ala Gly  Ile Trp Arg Val Glu  Cys Leu Ile Gly Glu  His Leu His
    1100                1105                1110

Ala Gly  Met Ser Thr Leu Phe  Leu Val Tyr Ser Asn  Lys Cys Gln
    1115                1120                1125

Thr Pro  Leu Gly Met Ala Ser  Gly His Ile Arg Asp  Phe Gln Ile
    1130                1135                1140

Thr Ala  Ser Gly Gln Tyr Gly  Gln Trp Ala Pro Lys  Leu Ala Arg
    1145                1150                1155

Leu His  Tyr Ser Gly Ser Ile  Asn Ala Trp Ser Thr  Lys Glu Pro
    1160                1165                1170

Phe Ser  Trp Ile Lys Val Asp  Leu Leu Ala Pro Met  Ile Ile His
    1175                1180                1185

Gly Ile  Lys Thr Gln Gly Ala  Arg Gln Lys Phe Ser  Ser Leu Tyr
    1190                1195                1200

Ile Ser  Gln Phe Ile Ile Met  Tyr Ser Leu Asp Gly  Lys Lys Trp
    1205                1210                1215

Gln Thr  Tyr Arg Gly Asn Ser  Thr Gly Thr Leu Met  Val Phe Phe
    1220                1225                1230

Gly Asn  Val Asp Ser Ser Gly  Ile Lys His Asn Ile  Phe Asn Pro
    1235                1240                1245

Pro Ile  Ile Ala Arg Tyr Ile  Arg Leu His Pro Thr  His Tyr Ser
    1250                1255                1260

Ile Arg  Ser Thr Leu Arg Met  Glu Leu Met Gly Cys  Asp Leu Asn
    1265                1270                1275

Ser Cys  Ser Met Pro Leu Gly  Met Glu Ser Lys Ala  Ile Ser Asp
    1280                1285                1290
```

Ala Gln Ile Thr Ala Ser Ser Tyr Phe Thr Asn Met Phe Ala Thr
1295 1300 1305

Trp Ser Pro Ser Lys Ala Arg Leu His Leu Gln Gly Arg Ser Asn
1310 1315 1320

Ala Trp Arg Pro Gln Val Asn Asn Pro Lys Glu Trp Leu Gln Val
1325 1330 1335

Asp Phe Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly
1340 1345 1350

Val Lys Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile
1355 1360 1365

Ser Ser Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn
1370 1375 1380

Gly Lys Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro
1385 1390 1395

Val Val Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg
1400 1405 1410

Ile His Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu
1415 1420 1425

Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr
1430 1435

<210> 1596
<211> 864
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1596
Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
1 5 10 15

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
20 25 30

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
35 40 45

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
50 55 60

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
65 70 75 80

```
Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
                85                  90                  95

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
            100                 105                 110

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
        115                 120                 125

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    130                 135                 140

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
145                 150                 155                 160

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
                165                 170                 175

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
            180                 185                 190

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
        195                 200                 205

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
    210                 215                 220

Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
225                 230                 235                 240

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro
                245                 250                 255

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
            260                 265                 270

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
        275                 280                 285

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
    290                 295                 300

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
305                 310                 315                 320

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                325                 330                 335

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
            340                 345                 350
```

```
Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
        355                 360                 365

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    370                 375                 380

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
385                 390                 395                 400

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
                405                 410                 415

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
            420                 425                 430

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
        435                 440                 445

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
    450                 455                 460

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
465                 470                 475                 480

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
                485                 490                 495

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
            500                 505                 510

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
        515                 520                 525

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
    530                 535                 540

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
545                 550                 555                 560

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                565                 570                 575

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
            580                 585                 590

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
        595                 600                 605

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
    610                 615                 620
```

```
Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
625                 630                 635                 640

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
                645                 650                 655

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            660                 665                 670

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
        675                 680                 685

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
    690                 695                 700

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
705                 710                 715                 720

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
                725                 730                 735

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
            740                 745                 750

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
        755                 760                 765

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
    770                 775                 780

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
785                 790                 795                 800

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
                805                 810                 815

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
            820                 825                 830

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
        835                 840                 845

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    850                 855                 860
```

<210> 1597
<211> 576
<212> PRT
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1597

```
Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser
1               5                  10                  15

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
            20                  25                  30

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
        35                  40                  45

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
    50                  55                  60

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
65                  70                  75                  80

Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
                85                  90                  95

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
            100                 105                 110

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser
        115                 120                 125

Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
    130                 135                 140

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
145                 150                 155                 160

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr
                165                 170                 175

Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
            180                 185                 190

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
        195                 200                 205

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly
    210                 215                 220

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
225                 230                 235                 240

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
                245                 250                 255
```

```
Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
            260                 265                 270

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
        275                 280                 285

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
    290                 295                 300

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser
305                 310                 315                 320

Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
                325                 330                 335

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser
            340                 345                 350

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
        355                 360                 365

Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
    370                 375                 380

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro
385                 390                 395                 400

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
                405                 410                 415

Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
            420                 425                 430

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro
        435                 440                 445

Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly
    450                 455                 460

Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
465                 470                 475                 480

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser
                485                 490                 495

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser
            500                 505                 510

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
        515                 520                 525
```

Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
530 535 540

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
545 550 555 560

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
565 570 575

<210> 1598
<211> 864
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1598
Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
1 5 10 15

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
20 25 30

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
35 40 45

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
50 55 60

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
65 70 75 80

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
85 90 95

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
100 105 110

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
115 120 125

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
130 135 140

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
145 150 155 160

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
165 170 175

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
180 185 190

```
Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
        195                 200                 205

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
    210                 215                 220

Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
225                 230                 235                 240

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro
                245                 250                 255

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
            260                 265                 270

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
        275                 280                 285

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
    290                 295                 300

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
305                 310                 315                 320

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                325                 330                 335

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
            340                 345                 350

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
        355                 360                 365

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    370                 375                 380

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
385                 390                 395                 400

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
                405                 410                 415

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
            420                 425                 430

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
        435                 440                 445

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
    450                 455                 460
```

```
Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
465                 470                 475                 480

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
                485                 490                 495

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
            500                 505                 510

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
        515                 520                 525

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
    530                 535                 540

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
545                 550                 555                 560

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                565                 570                 575

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
            580                 585                 590

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
        595                 600                 605

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
    610                 615                 620

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
625                 630                 635                 640

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
                645                 650                 655

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            660                 665                 670

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
        675                 680                 685

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
    690                 695                 700

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
705                 710                 715                 720

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
                725                 730                 735
```

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
740 745 750

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
755 760 765

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
770 775 780

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
785 790 795 800

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
805 810 815

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
820 825 830

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
835 840 845

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
850 855 860

<210> 1599
<211> 288
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1599
Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
1 5 10 15

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
20 25 30

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
35 40 45

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser
50 55 60

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
65 70 75 80

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
85 90 95

```
Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser
            100                 105                 110

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser
        115                 120                 125

Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser
    130                 135                 140

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser
145                 150                 155                 160

Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser
                165                 170                 175

Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
            180                 185                 190

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser
        195                 200                 205

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly
    210                 215                 220

Ala Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly
225                 230                 235                 240

Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser
                245                 250                 255

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr
            260                 265                 270

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
        275                 280                 285


<210> 1600
<211> 864
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
      polypeptide

<400> 1600
Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Pro
1               5                   10                  15

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
            20                  25                  30

Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
        35                  40                  45
```

```
Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
    50                  55                  60

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
65                  70                  75                  80

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
                85                  90                  95

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly
            100                 105                 110

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
        115                 120                 125

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    130                 135                 140

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
145                 150                 155                 160

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
                165                 170                 175

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
            180                 185                 190

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
        195                 200                 205

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro Ser Ala Ser Thr
    210                 215                 220

Gly Thr Gly Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
225                 230                 235                 240

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ala Ser Pro
                245                 250                 255

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
            260                 265                 270

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
        275                 280                 285

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro
    290                 295                 300

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
305                 310                 315                 320
```

```
Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                325                 330                 335

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly
            340                 345                 350

Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
        355                 360                 365

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    370                 375                 380

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ser Ser Thr
385                 390                 395                 400

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
                405                 410                 415

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
            420                 425                 430

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
        435                 440                 445

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
    450                 455                 460

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
465                 470                 475                 480

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
                485                 490                 495

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
            500                 505                 510

Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
        515                 520                 525

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
    530                 535                 540

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
545                 550                 555                 560

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
                565                 570                 575

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
            580                 585                 590
```

```
Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
        595                 600                 605

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
    610                 615                 620

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
625                 630                 635                 640

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
                645                 650                 655

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
            660                 665                 670

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
        675                 680                 685

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
    690                 695                 700

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
705                 710                 715                 720

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser Pro
                725                 730                 735

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
            740                 745                 750

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
        755                 760                 765

Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Pro
    770                 775                 780

Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro Gly Thr Ser Ser
785                 790                 795                 800

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
                805                 810                 815

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
            820                 825                 830

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
        835                 840                 845

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    850                 855                 860
```

<210> 1601
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1601
Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ser Ser
1 5 10 15

Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Thr Pro Gly Ser Gly Thr
20 25 30

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
35 40 45

Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser
50 55 60

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr
65 70 75 80

Ala Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser
85 90 95

Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ala Ser
100 105 110

Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser
115 120 125

Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser
130 135 140

<210> 1602
<211> 864
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1602
Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
35 40 45

```
Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
    50                  55                  60

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
65                  70                  75                  80

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
                85                  90                  95

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
            100                 105                 110

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
        115                 120                 125

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    130                 135                 140

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
145                 150                 155                 160

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
                165                 170                 175

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            180                 185                 190

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
        195                 200                 205

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
    210                 215                 220

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
225                 230                 235                 240

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
                245                 250                 255

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
            260                 265                 270

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        275                 280                 285

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
    290                 295                 300

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
305                 310                 315                 320
```

```
Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
                325                 330                 335

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
            340                 345                 350

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
        355                 360                 365

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    370                 375                 380

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
385                 390                 395                 400

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
                405                 410                 415

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            420                 425                 430

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
        435                 440                 445

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
    450                 455                 460

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
465                 470                 475                 480

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
                485                 490                 495

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
            500                 505                 510

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
        515                 520                 525

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala
    530                 535                 540

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
545                 550                 555                 560

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
                565                 570                 575

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
            580                 585                 590
```

```
Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
        595                 600                 605

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
    610                 615                 620

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
625                 630                 635                 640

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
                645                 650                 655

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
            660                 665                 670

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
        675                 680                 685

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
    690                 695                 700

Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
705                 710                 715                 720

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
                725                 730                 735

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
            740                 745                 750

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        755                 760                 765

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
    770                 775                 780

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
785                 790                 795                 800

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
                805                 810                 815

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
            820                 825                 830

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
        835                 840                 845

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    850                 855                 860
```

<210> 1603
<211> 576
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1603
Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
35 40 45

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
50 55 60

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
65 70 75 80

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
85 90 95

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
100 105 110

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
115 120 125

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
130 135 140

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
145 150 155 160

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
165 170 175

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
180 185 190

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
195 200 205

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
210 215 220

```
Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
225                 230                 235                 240

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
                245                 250                 255

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
            260                 265                 270

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        275                 280                 285

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
    290                 295                 300

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
305                 310                 315                 320

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
                325                 330                 335

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
            340                 345                 350

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
        355                 360                 365

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    370                 375                 380

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
385                 390                 395                 400

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
                405                 410                 415

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            420                 425                 430

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
        435                 440                 445

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
    450                 455                 460

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
465                 470                 475                 480

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
                485                 490                 495
```

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
500 505 510

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
515 520 525

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala
530 535 540

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
545 550 555 560

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
565 570 575

<210> 1604
<211> 864
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1604
Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
1 5 10 15

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
20 25 30

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
35 40 45

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
50 55 60

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
65 70 75 80

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
85 90 95

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
100 105 110

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
115 120 125

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
130 135 140

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
145 150 155 160

```
Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
                165                 170                 175

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            180                 185                 190

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
        195                 200                 205

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
    210                 215                 220

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
225                 230                 235                 240

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
                245                 250                 255

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
            260                 265                 270

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        275                 280                 285

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
    290                 295                 300

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
305                 310                 315                 320

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
                325                 330                 335

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
            340                 345                 350

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
        355                 360                 365

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    370                 375                 380

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
385                 390                 395                 400

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
                405                 410                 415

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            420                 425                 430
```

```
Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
        435                 440                 445

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
    450                 455                 460

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
465                 470                 475                 480

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
                485                 490                 495

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
            500                 505                 510

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
        515                 520                 525

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala
    530                 535                 540

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
545                 550                 555                 560

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
                565                 570                 575

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
            580                 585                 590

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
        595                 600                 605

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
    610                 615                 620

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
625                 630                 635                 640

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
                645                 650                 655

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
            660                 665                 670

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
        675                 680                 685

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
    690                 695                 700
```

```
Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
705                 710                 715                 720

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
                725                 730                 735

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
            740                 745                 750

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        755                 760                 765

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
    770                 775                 780

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
785                 790                 795                 800

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
                805                 810                 815

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
            820                 825                 830

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
        835                 840                 845

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    850                 855                 860


<210> 1605
<211> 288
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
      polypeptide

<400> 1605
Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
1               5                   10                  15

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
            20                  25                  30

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
        35                  40                  45

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
    50                  55                  60
```

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
65 70 75 80

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
85 90 95

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
100 105 110

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
115 120 125

Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
130 135 140

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
145 150 155 160

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
165 170 175

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
180 185 190

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
195 200 205

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
210 215 220

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
225 230 235 240

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
245 250 255

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
260 265 270

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
275 280 285

<210> 1606
<211> 864
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1606
Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
1 5 10 15

```
Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu
            20                  25                  30

Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
        35                  40                  45

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
    50                  55                  60

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
65                  70                  75                  80

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
                85                  90                  95

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala
            100                 105                 110

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
        115                 120                 125

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    130                 135                 140

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala
145                 150                 155                 160

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu
                165                 170                 175

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            180                 185                 190

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
        195                 200                 205

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
    210                 215                 220

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
225                 230                 235                 240

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
                245                 250                 255

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
            260                 265                 270

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        275                 280                 285
```

```
Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu
    290                 295                 300

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly
305                 310                 315                 320

Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
                325                 330                 335

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
            340                 345                 350

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu
        355                 360                 365

Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
    370                 375                 380

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr
385                 390                 395                 400

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
                405                 410                 415

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
            420                 425                 430

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
        435                 440                 445

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
    450                 455                 460

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
465                 470                 475                 480

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
                485                 490                 495

Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro
            500                 505                 510

Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
        515                 520                 525

Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala
    530                 535                 540

Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro
545                 550                 555                 560
```

```
Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
                565                 570                 575

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
            580                 585                 590

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
        595                 600                 605

Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro
    610                 615                 620

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr
625                 630                 635                 640

Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr
                645                 650                 655

Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
            660                 665                 670

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu
        675                 680                 685

Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr
    690                 695                 700

Ser Thr Glu Glu Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
705                 710                 715                 720

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu
                725                 730                 735

Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro
            740                 745                 750

Glu Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro
        755                 760                 765

Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Glu Pro
    770                 775                 780

Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly Ser Pro Thr
785                 790                 795                 800

Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
                805                 810                 815

Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Glu Pro
            820                 825                 830
```

Ala Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro
835 840 845

Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro
850 855 860

<210> 1607
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1607
Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
1 5 10 15

Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
20 25 30

Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
35 40 45

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
50 55 60

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
65 70 75 80

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
85 90 95

Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly Ser Pro Ala Gly
100 105 110

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
115 120 125

Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
130 135 140

<210> 1608
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1608
Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly
1 5 10 15

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly
20 25 30

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
35 40 45

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
50 55 60

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
65 70 75 80

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
85 90 95

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
100 105 110

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
115 120 125

Ser Gly Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
130 135 140

<210> 1609
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1609
Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser
1 5 10 15

Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser
20 25 30

Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly
35 40 45

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
50 55 60

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly
65 70 75 80

Ser Ala Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
85 90 95

Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Thr Glu
100 105 110

Pro Ser Glu Gly Ser Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser
115 120 125

Thr Glu Glu Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
130 135 140

<210> 1610
<211> 144
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1610
Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro Ala
1 5 10 15

Thr Ser Gly Ser Glu Thr Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
20 25 30

Ser Gly Pro Gly Ser Glu Pro Ala Thr Ser Gly Ser Glu Thr Pro Gly
35 40 45

Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Thr Ser Thr Glu
50 55 60

Pro Ser Glu Gly Ser Ala Pro Gly Thr Ser Glu Ser Ala Thr Pro Glu
65 70 75 80

Ser Gly Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly
85 90 95

Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu Gly Ser Pro Ala Gly
100 105 110

Ser Pro Thr Ser Thr Glu Glu Gly Thr Ser Glu Ser Ala Thr Pro Glu
115 120 125

Ser Gly Pro Gly Thr Ser Thr Glu Pro Ser Glu Gly Ser Ala Pro Gly
130 135 140

<210> 1611
<211> 19
<212> PRT
<213> Homo sapiens

<400> 1611
Met Gln Ile Glu Leu Ser Thr Cys Phe Phe Leu Cys Leu Leu Arg Phe
1 5 10 15

Cys Phe Ser

<210> 1612
<211> 6
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 1612
Gly Pro Glu Gly Pro Ser
1 5

<210> 1613
<211> 57
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 1613
atgcaaatag agctctccac ctgcttcttt ctgtgccttt tgcgattctg ctttagt 57

<210> 1614
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 1614
Ser Phe Ser Gln Asn Pro Pro Val Leu Lys Arg His Gln Arg
1 5 10

<210> 1615
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 1615
Gln Asn Pro Pro Val Leu Lys Arg His Gln Arg
1 5 10

<210> 1616
<211> 5
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 1616
Ser Phe Ser Gln Asn
1 5

<210> 1617
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 1617
Pro Pro Val Leu Lys Arg His Gln Arg
1 5

<210> 1618
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 1618
Leu Thr Pro Arg Ser Leu Leu Val
1 5

<210> 1619
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 1619
aggtgaatct ccdggtggyt cyagcggttc ygartc 36

<210> 1620
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 1620
acctgaytcr gaaccgctrg arccacchgg agattc 36

<210> 1621
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 1621
aggtagcgaa ggttcttcyg gtccdggyga rtcytc 36

<210> 1622
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 1622
acctgargay tcrcchggac crgaagaacc ttcgct 36

<210> 1623
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 1623
aggttcytcy gaaagcggtt cttcygargg yggtcc 36

<210> 1624
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 1624
acctggaccr ccytcrgaag aaccgctttc rgarga 36

<210> 1625
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 1625
aggttcyggt ggygaaccdt cygartctgg tagctc 36

<210> 1626
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 1626
aggttcgtct tcactcgagg gtac 24

<210> 1627
<211> 16
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide

<400> 1627
cctcgagtga agacga 16

<210> 1628
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide

<400> 1628
aggtagcccd gcwggytctc cdacytcyac ygarga 36

<210> 1629
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide

<400> 1629
accttcytcr gtrgargthg gagarccwgc hgggct 36

<210> 1630
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide

<400> 1630
aggtagcgaa cckgcwacyt cyggytctga racycc 36

<210> 1631
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide

<400> 1631
acctggrgty tcagarccrg argtwgcmgg ttcgct 36

<210> 1632
<211> 36

<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 1632
aggtacytct gaaagcgcwa cycckgartc yggycc 36

<210> 1633
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 1633
acctggrccr gaytcmggrg twgcgctttc agargt 36

<210> 1634
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 1634
aggtacytct acygaacckt cygarggyag cgcwcc 36

<210> 1635
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 1635
acctggwgcg ctrccytcrg amggttcrgt agargt 36

<210> 1636
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 1636
aggttctacy agcgaatcyc cktctggyac ygcwcc 36

<210> 1637
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide

<400> 1637
acctggwgcr gtrccagamg grgattcgct rgtaga 36

<210> 1638
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide

<400> 1638
aggtacytct acycckgaaa gcggytcygc wtctcc 36

<210> 1639
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide

<400> 1639
acctggagaw gcrgarccgc tttcmggrgt agargt 36

<210> 1640
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide

<400> 1640
aggtacytcy cckagcggyg aatcttctac ygcwcc 36

<210> 1641
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide

<400> 1641
acctggwgcr gtagaagatt crccgctmgg rgargt 36

<210> 1642
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide

<400> 1642
aggttcyacy agctctacyg cwgaatctcc kggycc 36

<210> 1643
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 1643
acctggrccm ggagattcwg crgtagagct rgtrga 36

<210> 1644
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 1644
aggtacycck ggyagcggta cygcwtcttc ytctcc 36

<210> 1645
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 1645
acctggagar gaagawgcrg taccgctrcc mggrgt 36

<210> 1646
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 1646
aggtagctct acyccktctg gtgcwacygg ytcycc 36

<210> 1647
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 1647
acctggrgar ccrgtwgcac cagamggrgt agagct 36

<210> 1648
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide

<400> 1648
aggttctagc ccktctgcwt cyacyggtac yggycc 36

<210> 1649
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide

<400> 1649
acctggrccr gtaccrgtrg awgcagamgg gctaga 36

<210> 1650
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide

<400> 1650
aggtgcwtcy cckggyacya gctctacygg ttctcc 36

<210> 1651
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide

<400> 1651
acctggagaa ccrgtagagc trgtrccmgg rgawgc 36

<210> 1652
<211> 57
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide

<400> 1652
aggtgcaagc gcaagcggcg cgccaagcac gggaggttcg tcttcactcg agggtac 57

<210> 1653

<211> 49
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 1653
cctcgagtga agacgaacct cccgtgcttg gcgcgccgct tgcgcttgc 49

<210> 1654
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 1654
Gly Ala Ser Ala Ser Gly Ala Pro Ser Thr Gly
1 5 10

<210> 1655
<211> 57
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 1655
aggtccagaa ccaacggggc cggccccaag cggaggttcg tcttcactcg agggtac 57

<210> 1656
<211> 49
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 1656
cctcgagtga agacgaacct ccgcttgggg ccggccccgt tggttctgg 49

<210> 1657
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 1657
Gly Pro Glu Pro Thr Gly Pro Ala Pro Ser Gly
1 5 10

<210> 1658
<211> 43

<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic primer

<400> 1658
tattcccgta cggccgccac catgcaaata gagctctcca cct 43

<210> 1659
<211> 32
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic primer

<400> 1659
ggtgacctcg agcgtagagg tcctgtgcct cg 32

<210> 1660
<211> 6
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 1660
Ala Arg Gly His Pro Phe
1 5

<210> 1661
<211> 5
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 1661
Asn Met His Thr Gly
1 5

<210> 1662
<211> 5
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 1662
Ser Pro Ala Thr Gly
1 5

<210> 1663
<211> 21

<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic primer

<400> 1663
gtgaactctc tagacccacc g 21

<210> 1664
<211> 37
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic primer

<400> 1664
ctcctcgagg tcgactcagt agaggtcctg tgcctcg 37

<210> 1665
<211> 92
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic primer

<400> 1665
cgaaagcgct acgcctgaga gtggccctgg ctctgagcca gccacctccg gctctgaaac 60

ccctgcctcg agcccaccag tcttgaaacg cc 92

<210> 1666
<211> 42
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic primer

<400> 1666
tgatatggta tcatcataat cgatttcctc ttgatctgac tg 42

<210> 1667
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic primer

<400> 1667
agcttgagga tccagagttc 20

<210> 1668
<211> 95
<212> DNA

<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
primer

<400> 1668
tctcaggcgt agcgctttcg cttgtcccct cttctgttga ggtgggggag ccagcaggag 60

aacctggcgc gccgttttga gagaagcttc ttggt 95

<210> 1669
<211> 39
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
primer

<400> 1669
tctcaaaacg gcgcgccagg tacctcagag tctgctacc 39

<210> 1670
<211> 30
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
primer

<400> 1670
tggtgggctc gaggctggcg cactgccttc 30

<210> 1671
<211> 39
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
primer

<400> 1671
tctcaaaacg gcgcgccagg tacctcagag tctgctacc 39

<210> 1672
<211> 30
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
primer

<400> 1672
tggtgggctc gaggctggcg cactgccttc 30

<210> 1673
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
primer

<400> 1673
ggggccgaaa cggccggtac ctcagagtct gctacc 36

<210> 1674
<211> 35
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
primer

<400> 1674
tgttcggccg tttcggcccc tggcgcactg ccttc 35

<210> 1675
<211> 29
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
primer

<400> 1675
ggggccgaaa cggccccggg agcgtcacc 29

<210> 1676
<211> 35
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
primer

<400> 1676
tgttcggccg tttcggcccc tgacccggtt gcccc 35

<210> 1677
<211> 55
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
primer

<400> 1677
caggacctct acggcgcgcc agcctcgagc gaacaaaaac tcatctcaga agagg 55

<210> 1678
<211> 55
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic

primer

<400> 1678
cctcttctga gatgagtttt tgttcgctcg aggctggcgc gccgtagagg tcctg 55

<210> 1679
<211> 70
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 1679
ctagcatgca aatagagctc tccacctgct tctttctgtg ccttttgcga ttctgcttta 60

gtgggtctcc 70

<210> 1680
<211> 70
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 1680
acctggagac ccactaaagc agaatcgcaa aaggcacaga aagaagcagg tggagagctc 60

tatttgcatg 70

<210> 1681
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic primer

<400> 1681
atgatggcat ggaagcctat 20

<210> 1682
<211> 91
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic primer

<400> 1682
atccctcacc ttcgccagaa ccttcagaac cctcaccttc agaaccttca ccagaacctt 60

caccatcttc cgcttcttca ttatttttca t 91

<210> 1683
<211> 93
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
primer

<400> 1683
ttctggcgaa ggtgagggat ctgaaggcgg ttctgaaggt gaaggtggct ctgagggttc 60

cgaatatgat gatgatctta ctgattctga aat 93

<210> 1684
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
primer

<400> 1684
tattctctgt gaggtaccag c 21

<210> 1685
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
primer

<400> 1685
tattctctgt gaggtaccag c 21

<210> 1686
<211> 67
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
primer

<400> 1686
gaagctggta cctcacagag aatatacaac gctttctccc caatccaggt gaaggttctg 60

gtgaagg 67

<210> 1687
<211> 47
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
primer

<400> 1687
aactctggat cctcaagctg cactccagct tcggaaccct cagagcc 47

<210> 1688
<211> 7
<212> PRT

<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 1688
Lys Leu Thr Arg Ala Glu Thr
1 5

<210> 1689
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 1689
Leu Gly Pro Val Ser Gly Val Pro
1 5

<210> 1690
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 1690
Gly Pro Ala Gly Leu Gly Gly Ala
1 5

<210> 1691
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 1691
Gly Pro Ala Gly Leu Arg Gly Ala
1 5

<210> 1692
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 1692
Ala Pro Leu Gly Leu Arg Leu Arg
1 5

<210> 1693

<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 1693
Pro Ala Leu Pro Leu Val Ala Gln
1 5

<210> 1694
<211> 4
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 1694
Gly Gly Ser Gly
1

<210> 1695
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 1695
Phe Asp Lys Leu Pro Arg Thr Ser Gly
1 5

<210> 1696
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 1696
Phe Lys Leu Met Val Trp Tyr
1 5

<210> 1697
<211> 44
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<220>
<221> MOD_RES
<222> (1)..(1)

<223> Any amino acid

<220>
<221> MOD_RES
<222> (44)..(44)
<223> Any amino acid

<220>
<223> see specification as filed for detailed description of substitutions and preferred embodiments

<400> 1697
Xaa Gly Ala Pro Gly Ser Pro Ala Gly Ser Pro Thr Ser Thr Glu Glu
1 5 10 15

Gly Thr Ser Glu Ser Ala Thr Pro Glu Ser Gly Pro Gly Ser Glu Pro
20 25 30

Ala Thr Ser Gly Ser Glu Thr Pro Ala Ser Ser Xaa
35 40

<210> 1698
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 1698
Arg His Gln Arg Glu Ile Thr Arg
1 5

<210> 1699
<211> 288
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 1699
Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
1 5 10 15

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
20 25 30

Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
35 40 45

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
50 55 60

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
65 70 75 80

```
Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro
                85                  90                  95

Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro
            100                 105                 110

Gly Thr Ser Ser Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser
        115                 120                 125

Thr Gly Ser Pro Gly Ala Ser Pro Gly Thr Ser Ser Thr Gly Ser Pro
    130                 135                 140

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
145                 150                 155                 160

Pro Ser Gly Ala Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
                165                 170                 175

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
            180                 185                 190

Gly Thr Pro Gly Ser Gly Thr Ala Ser Ser Ser Pro Gly Ser Ser Thr
        195                 200                 205

Pro Ser Gly Ala Thr Gly Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala
    210                 215                 220

Thr Gly Ser Pro Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro
225                 230                 235                 240

Gly Ser Ser Pro Ser Ala Ser Thr Gly Thr Gly Pro Gly Ala Ser Pro
                245                 250                 255

Gly Thr Ser Ser Thr Gly Ser Pro Gly Thr Pro Gly Ser Gly Thr Ala
            260                 265                 270

Ser Ser Ser Pro Gly Ser Ser Thr Pro Ser Gly Ala Thr Gly Ser Pro
        275                 280                 285


<210> 1700
<211> 6
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
      6xHis tag

<400> 1700
His His His His His His
1               5
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